STIC-Biotech/Ch mLib





From: Sent:

To:

Chan, Christina Tuesday, December 31, 2002 1:08 PM Holleran, Anne; STIC-Biotech/ChemLib RE: RUSH sequence search for 09/806,301

Subject:

Please rush. Thanks

Chris Chan TC 1600 New Hire Training Coordinator and SPE 1644 308-3973 CM-1, 9B19

Point of Contact: Mona Smith **Technical Information Specialist** CM1 6A01 Tel: 308-3278

-----Original Message-----

From: Holleran, Anne
Sent: Tuesday, December 31, 2002 1:06 PM
To: Chan, Christina

Subject:

RUSH sequence search for 09/806,301

Please approve and forward to STIC this RUSH sequence search request. This is an amendment due this biweek. Thansks.

Please search the following:

commercial and interference databases search of SEQ ID NO: 2(aa)

commercial and interference databases oligomer search of SEQ ID NO: 2(aa)

Anne Holleran AU: 1642 Tel: 308-8892 RM: 8e03

mailbox: 8e12

Searcher: H. Balon
Phone:
Location:
Date Picked Up: 1203
Date Completed: 1\3(03
Searcher Prep/Review:>_
Clerical:
Online time:

TYPE OF SEARCH:	
NA Sequences:	
AA Sequences:	
Structures:	
Bibliographic:	
Litigation:	
Full text:	
Patent Family:	
Other:	

/ENDOR/COST (where applic.)	
STN:	
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DRLink:	
Lexis/Nexis:	
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WWW/Internet:Other (specify):	
Other (5F5/)	

Point of Contact: Mona Smith Technical Information Specialism CM1 6A01 Tel: 308-3278

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Result
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Matches 51
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Zhao C., Nguyen T.X., Lehrer R.I.;
"Rabbit Lipophilins.";
submitted (SEP-2000) to the EMBL/Ge
EMBL; AF308614; AAG42802.1; -.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
SMART; SM0096; WTG; 1.
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O9GK67;
O1-MAR-2001 (TrEMBLrel. 16, Created)
O1-MAR-2001 (TrEMBLrel. 16, Last sequence update)
O1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Lipophilin AL.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                 SEQUENCE
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TISSUE-SUBMAXILLARY;

A Zhao C. Nguyen T.X., Lehrer R.I.;

Rabbit Lipophilins.";

L Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.

EMBL, AF308615; AAG42803.1;

PR InterPro; IPR000329; Uteroglobin.

DR Pfam; PF01099; Uteroglobin; 1.

DR Pfam; PF01099; Uteroglobin; 1.

DR YPRINTS; PR00486; UTEROGLOBIN.

DR SMART; SM00096; UTG; 1.

PARTICE 91 AA; 9723 MW; 579DCDDB813554C4 CRC64;
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O1-MAR-2001 (TIEMBLIEL 1
01-MAR-2001 (TIEMBLIEL 1
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InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
SEQUENCE 96 AA; 10510 MW; 48EC2972D0EA78C4 CRC
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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Cipophilin AL2.

Oryctolagus cuniculus (Rabbit).

Oryctolagus cuniculus (Rabbit).

Oryctolagus cuniculus (Rabbit).

Oryctolagus.
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01-MAR-2002 (TrEMBLrel.
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Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus
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TISSUE-LACRIMAL GLAND;
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                           GVKRCTDQMSLQKRSLIAEVLVKI-LKKCSV
                                                                                        MKILASILLISLALCCYEADAAACPTEIAESTANLLASESVERASLSKYGAPPEAVEAKL 60
                                                                                                                    MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
QVKRCTDKMSLGKRVLFGKVLGEIVLRTCTL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   47;
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                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            54.4%; Score 245; DE 52.8%; Pred. No. 1.16 tive 17; Mismatches
                                                                                                                                                                                                            51.4%; Score 231.5; DB 6 51.6%; Pred. No. 4.4e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16,
20,
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21,
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Last sequence update)
Last annotation updat
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Last sequence update)
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.le-21;
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A
                                                                                                                                                                                                                                DB 6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25;
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RESULT 6
Q8VD96
ID Q8VD96
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PRELIMINARI,
QBTD33;
Q1-JUN-2002 (TrEMBLrel. 21, Created)
Q1-JUN-2002 (TrEMBLrel. 21, Last sequence update)
Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE-LACRIMAL GLAND;
Zhao C., Nguyen T.X., Lehrer R.I.;
Rabbit Lipophilins.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF308616; AAG42804.1;
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
SEQUENCE 90 AA; 9943 MW; 410900DF7F3EB1BC CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.

Bingle C.D.;

"Human RYD5, a new secretoglobin.";

"Human RYD5, a new secretoglobin.";

Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases

EMBL; AY026938; AAK08972.1;

SEQUENCE 95 AA; 10457 MW; F629AF06C96D2392 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Ver
Mammalia; Eutheria; Lagomorpha; Leporidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-2001 (TrEMBLrel. 16, 01-MAR-2001 (TrEMBLrel. 16, 01-MAR-2002 (TrEMBLrel. 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9GK65;
01-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
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                                                                                                                                                             APPEAVAAKLGVKRCTDQMSLQKRSLIAEVLVKIL 85
                                                                                                                                                                                                                                                   MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFIS------
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                                                                                                                          VNEDAKAAMTELKSCRDGLQPMHKAELVKLLVQVL
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     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primates; Catarrhini; Hominidae;
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51.1%;
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                 Score 91; DB 4;
Pred. No. 0.0027;
6; Mismatches 3
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     PRT;
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Best Local
  Query Match
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Q8VFH6;
Q1-MAR-2002 (TrEMBLrel. 20, C
Q1-MAR-2002 (TrEMBLrel. 20, L
Q1-JUN-2002 (TrEMBLrel. 21, L
Q1factory receptor MOR114-B.
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01-MAR-2002 (TrEMBLrel. 2
01-MAR-2002 (TrEMBLrel. 2
01-JUN-2002 (TrEMBLrel. 2
CC10 protein precursor.
CC10.
                                            Receptor.
SEQUENCE
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SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cell 10 kDa protein.";
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases
EMBL; L37041; AAL31349.1; ...
InterPro; IPR003628; Uteroglbb_sub.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mesocricetus auratus
Eukaryota; Metazoa; C
Mammalia; Eutheria; F
                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gutierrez-Sagal R., Nieto A.; "Cloning and sequencing of a cDNA encoding hamster uteroglobin/clara
                                                                                                                                                    Submitted (JAN-2002) to the EMBL/GenBank/DDBJ EMBL; AY073550; AAL61213.1; -
InterPro; IPR000276; GPCR_Rhodpsn.
Pfam; PF00001; 7tm_1; 1.
                                                                                                                                                                                                                                                                                                                Zhang X., Firestein S.J.;
"The olfactory receptor gen
Nat. Neurosci. 0:0-0(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ProDom; PD012475; Uteroglbn_sub; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                             NCBI_TaxID-10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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                                                                                          PROSITE;
                                                                                                                                      PRINTS;
                                                                                                                                                                                                                                                 Adams M
                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PROSITE; PS00404; UTEROGLOBIN_2; UNKNOWN_1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MKIAITMAVVMLSVCCSSASSDTCPGFF-QVLEFLFMGSESSYEAALKFYNPGSDLQDSG
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                                                                                   PR00237; GPCRRHODOPSN.; PS00237; G_PROTEIN_RECEP_F1_1; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1
96 AA;
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                                              AΑ;
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10509
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Chordata; Craniata; Vertebrata; Euteleostomi;
Rodentia; Sciurognathi; Muridae; Cricetinae;
                                            35896 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18.9%;
  17.4%;
                                                                                                                                                                                                                                                                                                                                     gene
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21,
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Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 85; DB 11; Length 96; 
Pred. No. 0.014; 
7; Mismatches 36; Indels
                                                                                                                                                                                                                                                                                                                                       superfamily
Score 78.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         POTENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                         6C550A33DD277D24 CRC64;
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Length 317;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-FISCHER; TISSUE-OLFACTORY EPITHELIUM;
MEDLINE-92007724; PubMed-1915264;
Dear T.N., Boehm T., Keverne E.B., Rabbitts T.
"Novel genes for potential ligand-binding prot
the olfactory mucosa.";
EMBO J. 10:2813-2819(1991).
EMBL; X60661; CAA43068.1; -.
Interero; IPR000329; Uteroglobin.
SMART; SM0096; UTG; 1.
                                                                                                                                                                                                                                                                                                                                                                                                               Q05702
Q05702;
Q1-NOV-1996
Q1-NOV-1996
Q1-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Potential ligand-binding RYD5.
Adams M.;
Submitted (JAN-2002) to t
EMBL; AY073715; AAL61378
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rattus rattus (Black rat).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                   Zhang X., Firestein
"The olfactory recep
Nat. Neurosci. 0:0-0
                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                    NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                            Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRELIMINARY;
                                                                                                                                                y receptor gene of the control of th
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                                                                                                                                                                                                                                                                                                                                   Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17.2%;
28.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25.3%;
      the
8.1;
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Last sequ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 77.5; DB
Pred. No. 0.11;
15; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Last sequence up
         EMBL/GenBank/DDBJ
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3; Mismatches
                                                                                                                                                                                      superfamily
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25A4BBBB4977E247 CRC64;
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binding proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               320
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                                  databases
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; Murinae; Rat
                                                                                                                                                                                                                                                                                                                                   Euteleostomi; Murinae; Mus
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RA Arakawa T., Hara A., Fukunishi Y., Konno H., Adachi J., Fukuda S.,
RA Alzawa K., Izawa M., Nishi K., Kiyosawa H., Kondo S., Yamanaka I.,
RA Saito T., Okazaki Y., Gojobori T., Bono H., Kasukawa T., Saito R.,
RA Kadota K., Matsuda H.A., Ashburner M., Batalov S., Casavant T.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Fleischmann W., Gaasterland T., Gissi C., King B., Kochiwa H.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Schriml L.M., Staubli F., Suzuki R., Tomita M., Wagner L., Washio T.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Sakai K., Okido T., Furuno M., Aono H., Baldarelli R., Barsh G.,
RA Blake J., Boffelli D., Bojunga N., Carninci P., de Bonaldo M.F.,
RA Brownstein M.J., Bult C., Fletcher C., Fujita M., Gariboldi M.,
RA Lyons P., Marchionni L., Mashina J., Kamiya M., Lee N.H.,
RA Nordone P., Ring B., Ringwald M., Rodriguez I., Sakamoto N.,
RA Sasaki H., Toyo-oka K., Wang K.H., Weltz C., Whittaker C., Wilning L.,
RA Wunghar-Roris A. Yoshida W. Haeconawa V.
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Query Match
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Best Local Similarity
Matches 25; Conserv
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O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
MURINE CMAP (CYSTATIN F) (LEUKOCYSTATIN).
MURINE CMAP OR CST7.
                                  Pfam; PF00031; cystatin; 1.
SMART; SM00043; CY; 1.
SEQUENCE 167 AA; 18847 M
                                                                                                                "Functional annotation of a full-length mouse cDNA collection.";
Nature 409:685-690(2001).
EMBL; AB015224; BBA34940.1; -.
EMBL; AR004420; BAB23298.1; -.
EMBL; P01034; IG96.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRINTS; PR00237; GPCRRHODOPSN.
PROSITE; PS00237; G_PROTEIN_RECEP_F1_1;
PROSITE; PS50262; G_PROTEIN_RECEP_F1_2;
                                                                                                                                                                                                        Wynshaw-Boris A
Hayashizaki Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN-C57BL/6J; TISSUE-EMBRYO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-21085660; PubMed-11217851;
Kawai J., Shinagawa A., Shibata K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mus musculus (Mouse).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=10090;
                                                                              InterPro; IPR000010; Cystatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000276; GPCR_Rhodpsn.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rita M., Arakawa H., Yoshiuchi N.;
novel cystatin-like metastasis associated gene.";
mitted (JUN-1998) to the EMBL/GenBank/DDBJ databases
                                                                                                    MGI:1298217; Cst7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MDNKVCAILV---LCCWTSGLLVIIIPLGMILQLEFCD---SNTIDHFFCDASPLIKIS- 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MKLSVCLLLVTLALCCYQA------NAEFCPALVSELLDFFFI-SEPLFKLSL 46
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                                                                                                                                                                                                                        A., Yoshida K.,
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                                1.
18847 MW;
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25.3%;
 15.3%;
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Score 69; DB
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Pred. No. 2
                                                                                                                                                                                                                        Hasegawa Y., Kawaji H., Kohtsuki S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
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                                  61F776D8445095FE CRC64;
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RESULT 12
048907
ID 04890
AC 04890
DT 01-JU
DT 01-JU
DT 01-JU
DE MGATP
GN MRP2.
OC Eukar
OC Sperm

O48907 PRELIMINARY;
O48907;
O1-JUN-1998 (TIEMBLIEL 06, C
O1-JUN-1998 (TIEMBLIEL 06, L
O1-JUN-2002 (TIEMBLIEL 21, L
MGATP-energized glutathione S

Created)

PRT;

1623

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Last sequence update)
Last annotation update)
S-conjugate pump (EC 3.6.1

Arabidopsis thaliana (Mouse-ear cress). Eukaryota; Viridiplantae; Streptophyta; Embry Spermatophyta; Magnoliophyta; eudicotyledons;

Embryophyta;

core

eudicots;

Tracheophyta;

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104

PGLPPFEAFGLGVK 117

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RESULT
OAR 9018
ID 904
AC 904

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    048908;
048908;
01-JUN-1998
01-JUN-1998
01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lu Y.-P.,
Rea P.A.;
                                                                                                                                                                                                                                                SMARR; SMO0382; AAA, 1
PROSITE; PS00211; ABC_TRANSPORTER; 2.
ATP-binding; Hydrolase; Transport.
SEQUENCE 1622 AA; 182043 MW; 41AA
                                                                                                                                                                                                                                                                                                                                             InterPro; IPR003593; AAA_ATPase.
InterPro; IPR001140; ABCtranprtrTM.
InterPro; IPR003439; ABC_transportr.
pfam; pF00664; ABC_membrane; 2.
pfam; pF00005; ABC_tran; 2.
proDom; PD00006; ABC_transportr; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
Lu Y.-P., Li Z.-S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana (Mouse-ear cress).

Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosid.

eurosids II; Brassicales; Brassicacee; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MgATP-energized glutathione
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=3702;
  52
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  P - - PEAVAAKLGVK
                                                                                               LSVCLLLVTLALCCYQANAEFCPALVSELLDFFF-----ISEPLFKL----SLAKFDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GVKR---CTDQMSLQKRSLIAEVLVKILK
                                                   LVLCLYRIWLALKDHKVE-RFC--LRSRLYNYFLALLAAYATAEPLFRLIMGISVLDFDG
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                                                                                                                                                                             Similarity
                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Drozdowicz Y.M.,
                                                                                                                                                                          15.3%;
                                                                                                                                                                                                                                                  se; Transport.
182043 MW; 41AA158D694EA42A CRC64;
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Last annotation updat
S-conjugate pump (EC
                                                                                                                                                   9;
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                                                                                                                                                                          Score 69;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 2;
2; Mismatches
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                                                                                                                                                   Mismatches
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                                                                                                                                                                             DB
19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ssette transporter able chlorophyll catabolites
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pump (EC 3.6.1.3).
                                                                                                                                                                                                    10; Length 1622;
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                                                                                                                                                   Indels
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O64590 ID O6459 AC O6459 DT 01-AU DT 01-JU DT 01-JU DT AIC t GN ATABA AT
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Best Local :
                                                                                                                                                                                                                                                                               STRAIN-CV. COLUMBIA;
MEDLINE-20083487; PubMed-10617197;
Lin X., Kaul S., Rounsley S.D., Shea T.P., Benito M.-I., Town C.D.,
Pujii C.Y., Mason T.M., Bowman C.L., Barnstead M.E., Feldblyum T.V.,
Buell C.R., Ketchum K.A., Lee J.J., Ronning C.M., Koo H., Moffat K.9
Cronin L.A., Shen M., Vanaken S.E., Umayam L., Tailon L.J., Gill J.1
Adams M.D., Carrera A.J., Creasy T.H., Goodman H.M., Somerville C.R
Copenhaver G.P., Preuss D., Nierman W.C., White O., Eisen J.A.,
Salzberg S.L., Fraser C.M., Venter J.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AT2G34660.
Arabidopsis thaliana (Mouse-ear cress).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosideurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-AUG-1998 (TrEMBLrel. 07, 01-AUG-1998 (TrEMBLrel. 07, 01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transport glutathione S-conjugates and chlorophyll cat
functional comparisons with AtMRPI.";
Plant Cell 10:1-18(1998).
-!- SIMILARITY: BELONGS TO THE ABC TRANSPORTER FAMILY.
EMBL; AF020288; AAC04245.1; -.
HSSP; P13569; INBD.
                   Submitted (MAR-2000) to the EMBL/GenBank/DDBJ databases -!- SIMILARITY: BELONGS TO THE ABC TRANSPORTER FAMILY. EMBL; AC003096; AAC16268.1; -
                                                                                                                           SEQUENCE FROM N.A. STRAIN-CV. COLUMBIA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      064590;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ProDom; PD000006; ABC_transportr; 2.
SMART; SM00382; AAA; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABC transporter (AtMRP2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 064590
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATP-binding; Hydrolase; Transport SEQUENCE 1623 AA; 182073 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS00211; ABC_TRANSPORTER; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF000664; ABC_membrane; 2. Pfam; PF00005; ABC_tran; 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR001140; ABCtranprtrTM.
InterPro; IPR003439; ABC_transportr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
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                                                                                                          Lin X.;
                                                                                                                                                                                                                                                         Salzberg S.L., Fraser C.M., Venter J.C.;
"Sequence and analysis of chromosome 2 of the plant Arabidopsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=3702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LVLCLYRIWLALKDHKVE-RFC--LRSRLYNYFLALLAAYATAEPLFRLIMGISVLDFDG 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27;
                                                                                                                                                                                                             402:761-768(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Drozdowicz Y.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               182073 MW;
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Pred. No. 19;
9; Mismatches
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Last annotation updat
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19;
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RESULT
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Pfam; PF00005; ABC_tran; 2.
ProDom; PD000006; ABC_transportr; 2.
SMART; SM00382; AAA; 1.
PROSITE; PS00211; ABC_TRANSPORTER; 2
ATP-binding; Transport.
SEQUENCE 1623 AA; 182129 MW; F07.
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STRAIN-LANDSBERG ERECTA;
MEDLINE-98189246; PubMed-9528668;
MEDLINE-98189246; PubMed-9528668;
Marin E., Leonhardt N., Vavasseur A., Forestier C.;
"Cloning of AtmrP1, an Arabidopsis thaliana cDNA encoding of the mammalian multidrug resistance-associated protein."
Biochim. Biophys. Acta 1369:7-13(1998).
-i-SIMILARITY: BELONGS TO THE ABC TRANSPORTER FAMILY.
EMBL; AF014960; AAC49988.1; -.
HSSP; P13569; INBD.
                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00064; ABC_membrane; 2. Pfam; PF00005; ABC_tran; 2. ProDom; PD000006; ABC_transportr; SMART; SM00382; AAA; 1.
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;

Spermatophyta; Magnoliophyta; eudicotyta; edicots; Rosid

eurosids_II; Brassicales; Brassicaceae; Arabidopsis.
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Ol-JAN-1998 (TrEMBLrel. 05, Last sequence update)
Ol-JUN-2002 (TrEMBLrel. 21, Last annotation upda:
Multidrüg resistance-associated protein 2.
                                                                                                                                                                                                                                                                                                      PROSITE; PS00211; ABC_TRANSPORTER; ATP-binding; Transport. SEQUENCE 1623 AA; 181968 MW; 11
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InterPro; IPR003439; ABC_transportr.
                                                                                                                                                                                                                                                                                                                                                                                SMART; SM00382; AAA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR003593; AAA_ATPase.
InterPro; IPR001140; ABCtranprtrTM.
InterPro; IPR003439; ABC_transportr
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                                                                                                                                                   9 LVTLALCCYQ-----ANAEFCPALVSELLDFFF-----ISEPLI'KL----SLAKF
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DGPGLPPFEAFGLGVK 117
                                                DAP - - PEAVAAKLGVK
                                                                                                  LVLLVLCLYRIWLTLKDHKVERFC--LRSRLYNYFLALLAAYATAEPLFRLIMGISVLDF
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36.5%;
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35.5%;
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                                                                                                                                                                                                Pred. No. 22;
6; Mismatches
                                                                                                                                                                                                                             Score 68.5;
Pred. No. 22;
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RESULT 15
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DT 01-MA
DT 01-MA
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DE Putat
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RC STRAIN-NIPPONBARE;

RA MCCOmble W.R., de la Bastide M., Spiegel L., Preston R., Kirchoff K.,

RA MCCOmble W.R., de la Bastide M., Spiegel L., Preston R., Kirchoff K.,

RA MCCOmble W.R., de la Bastide M., Balija V., Bell M., Baker J.,

RA MCCOmble W.R., Mascimento L., Zutavern T., Balija V., Bell M., Baker J.,

RA Santos L., Miller B., Katzenberger F., Muller S., King L., Yang C.,

RA O'Shaughnessy A., Palmer L., Dedhia N.;

"Genomic sequence for Oryza sativa, Nipponbare strain, clone

"Genomic sequence for Oryza sativa, Nippon
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 20
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QBW3C2;
Q1-MAR-2002 (TrEMBLrel. 20, Created)
Q1-MAR-2002 (TrEMBLrel. 20, Last sequence update)
Q1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Putative polyprotein (Hypothetical 33.4 kDa protein).
QSJNBA0046L02.13 QR QJ1136E01.1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
Wing R.A., Yu Y., Soderlund C., Chen M., Kim H.-R., Rambo T.,
Saski C., Henry D., Oates R., Simmons J.;
"Rice Genomic Sequence.";
submitted (FEB-2002) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Oryza sativa (Rice).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
NCBI_TaxID=4530;
                                                                                                                                                                                                               / Match 15.1%; Score 68; DB 10; Length 290; Local Similarity 36.4%; Pred. No. 4.6; ndels 20; Conservative 8; Mismatches 23; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               4.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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Minimum
Maximum
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Post-processing: Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Searched:
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length: 2000000000
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90
              January
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SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1985.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1985.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1987.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1989.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1999.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1999.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1991.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1993.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1995.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1995.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1996.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1997.DAT:

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SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1999.DAT:

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SIDS2/gcgdata/geneseq/geneseqp-emb1/AA2000.DAT:

SIDS2/gcgdata/geneseq/geneseqp-emb1/AA2000.DAT:
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Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                       /SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1982.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1983.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-emb1/AA1984.DAT:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Search time 35 Seconds
(without alignments)
342.644 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

45	44	43	42	41	40	39	38	37	36	35	34	ω ω	32	31	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	13	12	11
15	15	15	15	15	15	15	15	15	15	22	22	37	39	45	45	48	50	50	50	50	50	52	52	52	52	62	66	69	69	79	90	90	90	90
16.7		16.7	•	•	16.7	16.7	16.7	16.7		•	٠	٠								٠			57.8				٠		76.7	•	•		100.0	100.0
16	16	16	15	15	15	15	15	15	15	22	22	90	90	45	45	88	50	50	50	50	50	90	90	53	53	90	90	69	69	89	120	117	90	90
19	19	19	21	21	21	21	21	19	19	21	19	22	22	21	19	22	23	22	22	22	22	22	22	21	21	22	22	21	19	23	20	22	23	22
AAW54276	AAW54274	AAW54273	AAB07506	AAB07505	AAB07504	AAB07503	AAB07502	AAW54275	AAW54272	AAB07507	AAW54277	AAE07526	AAE07522	AAB07508	AAW54278	AAE07523	ABG43285	AAM33628	AAM73426	AAM60754	ABB40007	AAE07527	AAE07524	AAY64680	AAG00474	AAE07521	AAE07525	AAB07509	AAW54279	AAO20554	AAY48606	ABB11907	96	AAB31681
		BU101 antigenic pe	Amino acid sequenc			Amino acid sequenc	O.	antigenic	anti	acid seque	ซ	_	lipo	acid seque	BU101 antigenic pe	lipophil	Human peptide enco	Peptide #7665 enco		Human brain expres	O	lipophilin	lipophili	-	Human secreted pro	Human lipophilin B	Human lipophilin B		antigenic p	in of hu	breast		man endometrial	An endometrial spe

ALIGNMENTS

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RESULT 1
AAW35803
ID AAW35803
XX AAW35803
XX AAW35803
XX Endc
Clark
KW Clark
KW Clark
KW Phoso
KW infl
KW phoso
KW infl
KW Phoso
XX Endc
X
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phospholipase A2 inhibitor; polychlorinatiad biphenyl; antiaggregant;
inflammation; asthma; rhinitis; cystic fibrosis; airway disease;
neoplasia; atopy; therapy; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW35803 standard; Protein; 90 AA.
                                                                                         21-MAR-1996;
                                                                                                                                                                              21-MAR-1996;
                                                                                                                                                                                                                                                                            25-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                 W09734997-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Endometrial specific steroid-binding factor II; ESF II; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human endometrial specific steroid-binding factor II.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW35803;
    (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                         96WO-US03857
                                                                                                                                                                                   96WO-US03857
                                                                                                                                                                                                                                                                                                                                                                                                                                        /label= Sig_peptide
22..90
/label= Mat_protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence comprises human endometrial specific steroid binding catchity, binds to polychlorinated biphenyl compounds, reduces foreign protein antigenicity, inhibits monocyte and neutrophil chemotaxis and phagocytosis, inhibits monocyte and neutrophil chemotaxis and phagocytosis, inhibits platelet aggregation, regulates eicosanoid levels in the human uterus and controls the growth of endometrial cells. The amino acid sequence was deduced from a cDNA clone (see AAM39831) derived from cyclohaximide-treated CEM cells. ESF I (see AAM39832) and ESF III see AAM35804) are also claimed. Human ESF II has about 49% identity with rat prostatic steroid-binding protein. Recombinant ESF I, and III can be expressed in host cells for use in claimed methods (a) for treating a patient in need of ESF I, II or III (including expression of the polypeptide in vivo) and (b) for identifying compounds which bind to and inhibit activation of the ESF polypeptide. hESF I, II and III may be used to treat inflammation, asthma, rhinitis, cystic fibrosis, airway disease, neoplasia and atopy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                            Human steroid-binding protein C1; hSBP1; hSBP2; breast cancer; prob gene therapy vector; ribozyme; probe; hybridisation; amplification; antibody; immunoassay.
                                                                                                                                                                                                                                                                 Amino acid sequence of the human steroid binding protein
                                                                                                                                                                                                                                                                                                                                                     AAW59776 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human endometrial specific steroid-binding factor I, used to treat inflammation, asthma, rhinitis, cystic
                Akerblom
                                        (INCY-) INCYTE PHARM INC
                                                                    12-NOV-1996;
                                                                                                07-NOV-1997;
                                                                                                                             22-MAY-1998
                                                                                                                                                         WO9821331-A1
                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                              12-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 18; Page 63-64; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     airway disease, neoplasia, atopy etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gentz RL,
                                                                                                                                                                                                                                                                                                                                                                                                                             61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
              IE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
             Goli SK,
                                                                    96US-0747547
                                                                                              97WO-US20674
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             Hawkins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 90; DB 18;
Pred. No. 2.2e-82;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                       B
             PR,
             Hillman
             JĽ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
             Murry LE;
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RESULT 3
AAW54271
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Best Local S
Matches 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        monitoring and treatment of breast cancer. HSBP1 and hSBP2 are useful as markers for breast cancer, i.e. measuring levels of hSBP1 and hSBP2 used for diagnosis or monitoring the disease, to identify subjects at risk and to discriminate between different forms of cancer for selection of appropriate therapies. They may also be used for drug screening. Nucleic acids encoding hSBP1 and hSBP2 can be used in gene therapy vectors to overexpress the steroid-binding proteins, preventing binding of steroids, or antisense sequences, ribozymes. Their nucleic acids can also be used for the diagnosis and monitoring (by quantifying expression of hSBP), as source of probes for hybridisation and amplification of genomic or related sequences for studying regulation of gene function and for mapping the genomic sequence. Antibodies are used as diagnostic reagents in standard immunoassays for hSBP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This
                                                         Billing-medel PA, Coh
Gordon J, Granadosen,
                                                                                                                                                                                                                                                                                       BU101 antigenic peptide epitope 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New human steroid binding diagnosis, monitoring and
                                                                                            (ABBO ) ABBOTT LAB.
                                                                                                                    15-AUG-1997;
19-AUG-1996;
                                                                                                                                                        19-AUG-1997;
                                                                                                                                                                                26-FEB-1998
                                                                                                                                                                                                        WO9807857-A1
                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                 BU101;
                                                                                                                                                                                                                                                                                                                29-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                         AAW54271;
                                                                                                                                                                                                                                                                                                                                                                 AAW54271 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C1 (hSBP1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1998-297935/26.
N-PSDB; AAV41579.
                                                                                                                                                                                                                                                      immunisation;
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                                                                                                                                                                                                                                                                breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; 70pp;
                                                                                                                                                                                                                                                    cancer;
drug scr
                                                                                                                    97US-0912276
96US-0697105
                                                                                                                                                        97WO-US14665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  no acid sequence of the human steroid-binding protein in the method of the invention for the diagnosis,
                                              Russell JC,
                                                                   Cohen M,
                                                                                                                                                                                                                                                                                                                                                                 Protein;
                                                                                                                                                                                                                                                      screening;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
                                                                                                                                                                                                                                                                diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                        en M, Colpitts TL,
Hodges SC, Klass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       proteins C1 and treating breast
                                                                                                                                                                                                                                                                                                                                                                 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 90; DB 19;
Pred. No. 2.2e-82;
                                                                                                                                                                                                                                                     epitope
                                                                                                                                                                                                                                                                                                                                                                 Ā
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                prevention; treatment; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C2 - useful cancer, and
                                                          MR,
                                                                      Friedman PN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                          Kratochvil JD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
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N-PSDB; AAV26461 WPI; 1998-169161/15 Roberts-rapp L,

Stroupe

SD;

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ARRESULT 4
AAW89613
AAW899613
AC AAW89
AC AAW8
AC AAW8
AC Endo
XX Endo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cells transformed with a recombinant expression system comprising a consequence derived from the BUJO1 open reading frame and with at least 50 percent identity to the sequences shown in AAV26458 to AAV26461 are used to produce BUJO1 polypeptides containing at least 1 epitope. These are used to detect BUJO1 specific antibodies which are used correspondingly to detect BUJO1 antigens. The BUJO1 polynucleotide sequences can be used in a method for detecting the presence of a target BUJO1 polynucleotide. The various assays are used for diagnosis, prognosis, staging, monitoring, treating and preventing diseases of the breast (especially cancer and its metastases), and also for determining susceptibility. The BUJO1 polypeptides are also useful in drug screening, e.g. to identify and appreventing diseases of the breast (especially cantagonists of BUJO1, potentially useful therapeutically and a targets for therapy. The antibodies are also useful for targeted drug delivery and therapeutically to neutralise BUJO1 polypeptides. Fragments of the BUJO1 nucleic acid are useful as probes and primers, e.g. for detection of altered gene expression or in fluorescent in situ hybridisation, also in gene therapy to generate antisense or ribozyme molecules or for generatic immunisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loc
Matches
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Treatment of endometrial cancer, mammary cancer, endometriosis or endometrial fibroids - comprises administering endometrial steroid
                                                                                                                                                                                                                                                                           09-JUN-1998;
                                                                                                                                                                                                                                                                                                                       17-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                          endometriosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endometrial steroid binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endometrial steroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAW89613;
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                                                                        N-PSDB; AAX00069
                                                                                                                                                                                  (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                                                                              09-JUN-1997;
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                                                                                           1999-080843/07.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      fibroid; mammary cancer.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             II; ESBPII; cancer; detection;
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RESULT 5
AAB13787
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for diagnosing the above mentioned diseases comprising analysing the abnormally high level of ESBPII polypeptice in cells, tissues and bodily fluids; and (2) a diagnostic method for the diseases described above comprising analysing the abnormally high or low transcription level of ESBPII in cells, tissues and bodily fluids. The methods can be used to diagnose, treat, and monitor the progression, remission or recurrence of abnormal cell growth, such as cancers, especially endometrial and mammary cancer, and endometriosis and endometrial fibroids. The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A method has been developed for the treatment of endometrial cancer, mammary cancer, endometriosis or endometrial fibroids. The method comprises administering endometrial steroid binding protein II (ESBPII) antagonist. Also described in the present invention are: (1) a method antagonist.
Multimeric polypeptide antigen and antikody specific useful for diagnosing, detecting and treating breast
                                                      WPI; 2000-442366/38.
N-PSDB; AAA64846.
                                                                                                                                                                                                                                                                                                                                                                      Key
                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; breast cancer; breast disease detection; mammaglobin; uteroglobin; BU101; endometrial; cytostaric.
                                                                                                       Colpitts TL,
                                                                                                                                                                                                            20-DEC-1999;
                                                                                                                                                                                                                                               22-JUN-2000
                                                                                                                                                                                                                                                                                                                                                      Misc-difference
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                                                                                                                                       (ABBO ) ABBOTT
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                                                                                                       Russell JE;
                                                                                                                                         LAB
                                                                                                                                                                            98US-0215818
                                                                                                                                                                                                              99WO-US30489
                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                      'label= Leu
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2.2e-82;
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   to the cancer
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Claim 1; Page 124; 124pp; English

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Query Match
Best Local
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                                                                                           The present sequence represents a BUI01 polypeptide. The BUI01 gene is transcribed from breast tissue. The specification describes a method detecting the presence of a target BUI01 polynuclectide in a test sample. The method comprises contacting the sample with at least one BUI01-specific polynuclectide (AAA58875-80), and detecting bound build sample with a sample
                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                                          Detecting presence of target BU101 polynucleotide in sample detection of breast cancer, comprises contacting sample with BU101-specific polynucleotide and determining binding -
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
                        polynucleotides. The method and BU1010 polynucleotides are useful for detecting the presence of BU101 polynucleotides. The methods may be used for the diagnosis of breast disease, indicated by the formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Billing-medel PA,
Granados EN, Hodo
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                       antiallergic, and cytostatic properties. The polynucleotides are used in companies the same used in the polynucleotides are used in the polynucleotides are used in the polynucleotides in vivo to treat and/or prevent inflammation, asthma, rhinitis, cystic fibrosis, air way disease, neoplasia and atopy. The polynucleotides are also used to inhibit phospholipase A2 activity, bind polychlorinated biphenyls, reduce foreign protein antigenicity, inhibit monocyte and neutrophil chemotaxis and phagocytosis, inhibit platelet aggregation, regulate eicosanoid levels in the human uterus and control the growth of endometrial cells. The polynucleotides are also useful for detecting complementary polynucleotides are used to detect complementary polynucleotides are dispossic reagent. Detection of a mutated form of hESF I, II and III associated with a dysfunction will provide a diagnostic tool that can define diagnosis of a disease or susceptibility to a disease which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Endometrial specific steroid-binding factor; human; asthma; rhinitis; cystic fibrosis; air way disease; eicosanoid level regulator; chemotaxis inhibitor; er
                                                                                                                                                                                                                                                    This invention relates to nucleic acid molecules encoding portions of the human endometrial specific steroid-binding factors I, II, and III. Also included in the invention are hESF I, II, and III polypeptide sequences. The nucleotide sequence exhibit antiasthmatic, antiinflammatory,
                                                                                                                                                                                                                                                                                                                                    Claim 1; Fig
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                                                                                                                                                                                                                                                                                                                                                                  gene encoding human endometrial specific steroid-binding factor and III which is useful for treating asthma, rhinitis, cystic sis, airway disease and neoplasia -
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Best Local :
                                                                    The present sequence represents a human endometrial specific steroid-binding protein (ESBP) II. The ESBPII protein is overexpressed in breast tumours. The specification describes a method for diagnosing prostate or a gynaecological cancer. The method comprises measuring levels of ESBPII in cells, tissues or body fluids of a patient, and comparing this to levels from a normal control, where a variance in levels indicates cancer. The method is used to diagnose, stage, monit image or treat prostate or gynaecological cancer. The gynaecological cancers include breast, endometrial, ovarian and uterine cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      endometrial cancer. They are also useful for chromosome identification. The present sequence represents a hESF II protein sequence identified in the invention.
                                                                                                                                                                                                                       Diagnosing, staging, monitoring, imaging and treating prostate and gynaecological cancers by measuring levels of endometrial specific steroid-binding protein (ESBP)II expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; endometrial specific steroid-binding protein II; ESBPII; breast tumour; prostate cancer; gynaecological cancer; cancer; endometrial cancer; ovarian cancer; uterine cancer.
                                                                                                                                                                                                 Claim 6; Page 31-32; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A human endometrial specific steroid-binding protein
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                                                          Query Match
                                                                                                                                                                     expressed in TI, and performing quantitative polymerase chain reaction (PCR) analysis on SP1 to identify second subset of (P). The method is useful for determining the presence or absence of a cancer cell in a patient, monitoring the progression of cancer in a patient using a biological sample such as blood, serum, lymph nodes, bone marrow, sputum, urine or a tumour biopsy sample. The methods are useful for determining the presence or or absence of or monitoring progression of prostate, breast, colon, overian, lung, head and neck, lymphoma, leukemia, melanoma, liver, gastric, kidney, bladder, pancreatic or endometrial cancer. The present sequence represents the lipophilin B pclypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying tissue (tumour)-specific polynucleotides overexpressed tissue of interest as compared to control tissue, for detecting car cells in patient, comprises DNA microarray analysis or quantitative polymerase chain reaction
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20-JUL-2000; 2000US-219862P.
27-JUL-2000; 2000US-221300P.
18-DEC-2000; 2000US-256592P.
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cancer; B726P; Lipopl
                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to identifying thissue-specific polynucleotides (P) that involves performing a genetic subtraction to identify pool of (P)
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DB; AAI67269.
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   h
Similarity 100
90; Conservative
                                                                                                                       90 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ction; DNA microarray analysis;
Lipophilin B; mammaglobin.
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                              .0%;
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   Score 90; DB
Pred. No 2.2
); Mismatches
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2.2e-82;
hes 0;
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                                                          Length
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hESFI; hESFII;

way disease;
chemotaxis;

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ARESULT 10
AREO7518
ID AREO7
XX AREO7
XX O6-NC
XX O6-NC
XX Cance
XX Human
XX Human
XX Homo
XX Homo
XX Homo
XX Homo
XX Homo
XX 11-FE
PR 28-JU
XX 12-FE
PR 18-FE
XX 11-FE
PR 19-FE
XX 11-FE
PR 11-
   RESULT 11
AAB31681
ID AAB31
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a complex comprising a lipophilin-like polypeptide linked by at least one disulphide bond to a second lipophilin-like polypeptide. Lipophilin-like protein are members of uteroglobin superfamily. Lipophilin-like proteins are useful in the preparation of vaccines. The complex containing lipophilin-like proteins are useful for treating or preventing breast, ovarian or prostate cancer. The complex is also used for determining the presence or absence of cancer in a patient, or monitor the progress of cancer in a patient. Lipophilin DNA is also useful in gene therapy. The present sequence is human lipophilin B protein.
                                           AAB31681 standard;
                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Page 72; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        prostate cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel isolated complex two least one disulfide bond, u
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28-JUN-2000;
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                                                                                                                                                                                                            GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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DB; AAD13756.
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                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                                                                                                                                         100.0%;
ilarity 100.0%;
Conservative 0
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2000US-0215735
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in B; cytostatic;
ovary; prostate.
                                         Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vallieve-Douglass J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        o lipophilin-like used to treat or
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                                                                                                                                                                                                                                                                                         Score 90; DB 22;
Pred. No. 2.2e-82;
; Mismatches 0;
                                           B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptides linked by at prevent breast, ovarian or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Houghton RL,
                                                                                                                                                                                                                                                                                                                              Length
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RESULT 12
ABB09634
ID ABB09
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AC ABB09
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DT 29-MA
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Best Local S
Matches 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New human endometrial specific steroid binding factors, treating and preventing inflammation, asthma, rhinitis, fibrosis, airway disease, neoplasia and atopy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Key
Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; endometrial specific steroid binding factor; hESF; hESFI; hESFIII; inflammation; asthma; rhinitis; cystic fibrosis; airway neoplasia; atopy; phospholipase A2; polychlorinated biphenyl; chephagocytosis; platelet aggregation; elcosanoid; endometrial cell.
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Fig 2; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB; AAF25213.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAR-1996;
21-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 An endometrial specific steroid binding
   29-MAY-2002
                                         ABB09634;
                                                                            ABB09634
                                                                                                                                                                                                                                                                                                                                                                                                                                            therapeutic purposes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-APR-2001
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                                                                                                                                                                                                                                                 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                            GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
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                                                                                                                                                                                                                                                                                                                            similarity 100.0%; 90; Conservative (
                                                                            standard; Protein;
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 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96US-0014724.
97US-0821451.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "signal peptide"
                                                                              90
                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                            Score 90; DB 22;
Pred. No. 2.2e-82;
); Mismatches 0;
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ABB11907
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Human
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Best Local (
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21-MAR-1997;
08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                  The present sequence represents a endometrial specific steroid-binding factor (hESF) II. The full length protein has a molecular weight of 9.9 kDa. The protein has homology to rat prostatic steroid-binding protein C2. Antibodies which bind hESF proteins, such as hESF I, hESF II, and hESF III are useful for isolating or to identify clones expressing the polypeptides or to purify the polypeptides by affinity chromatography. Agonstst and antagonists in hESF proteins are useful for treatment and antagonists of hESF proteins are useful
Human; cytokine; cell proliferation; cell differentiation; haematopolesis regulation; tissue growth; immunomodulator;
                                        Human breast tumour-associated protein homologue, SEQ ID NO: 2277.
                                                                   11-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antibody specific for human endometrial specific steroid-binding factor (hESF) III, useful for detecting hESF III protein in biological sample and to isolate or identify clones expressing the protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAY-2000;
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                                                                                                                      ABB11907 standard;
                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Fig 1; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN
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                                                                                                                                                                                                                                          -
                                                                                                                                                                                                                                                    1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                             GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                      MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                                                                                                                                      GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  endometrial specific steroid-binding factor (hESF)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            endometrial specific steroid-binding factor;
ic steroid-binding protein; hESF I; hESF II;
                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                 90
                                                                                                                                                                                                                                                                                           Conservative
                                                                  (first entry)
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97US-0821451.
99US-0263810.
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22..90
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                                                                                                                     peptide; 117
                                                                                                                                                                                                                                                                                                                                                                          preventing susceptibility to asthma.
                                                                                                                                                                                                                                                                                                         100
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Pred. No. 2.2e-82;
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  activin;
             growth factor;
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thrombolytic activities; receptor or ligand activities; or may be convolved in oncogenesis, cancer cell preliferation or metastasis. Compending on their biological activities, polypeptides and nucleotides of the invention are useful for preventing, treating or ameliorating medical conditions, e.g., by protein or gene the rapy. Such conditions include cancers, haematopoietic disorders (e.g., myeloid or lymphoid cell clasorders), chronic inflammatory conditions (e.g., asthma or arrhritis), proliferative retinopathy, atherosclerosis, coronary heart disease, arterial ischaemia, bone disorders (e.g., osteoporosis), and abnormal vascular growth. Polypeptides involved with tissue regeneration and repair (or nucleic acids encoding them) may be used to promote wound healing (e.g., of burns, inclsions and ulcers), while those with immunomodulatory activities may be used in the treatment of viral, bacterial and fungal infections in addition to immune disorders. Polypeptides with growth factor activity may be used in cell cultures to promote cell growth. For example, such polypeptides may be used to result can be used to augment or replace 'sells damaged by illness, that can be used in the diagnosis of the above conditions, and in drug soreening techniques. The polypeptides and nucleotides may also be used in the diagnosis of the above conditions, and in drug soreening techniques. The present sequence represents a novel human collaboration of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequences ABB10981-ABB12330 represent 13:0 novel human polypeptides, and sequences ABA08225-ABA09574 represent nu:leit acids encoding them. The invention also relates to vectors and recombinant host cells comprising a nucleotide of the invention, methods of producing the novel polypeptides, antibodies against the polypeptides, methods of dentifying compounds which polypeptides in a sample, and methods of identifying compounds which bind to polypeptides of the invention. Although novel, many of the polypeptides of the invention have homology to known proteins, thereby giving an insight into their probable biological activities, and hence potential therapeutic applications. The polypeptides of the invention may have various activities, including cytokine, cell proliferation or cell differentiation activities; stem cell growth factor activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inhibin; chemotaxis; chemokinesis; thrombolysis; oncogenesis; proliferation; metastasis; cancer; tumour; haematopoietic disorder myeloid cell disorder; proliferative retinopathy; chronic inflammatory condition; proliferative retinopathy; atherosclerosis; coronary heart disease; arterial ischaemia; bone disorder; osteoporosis; vascular growth disorder; tissue regeneration; wound healing; infection; immune disorder; cell culture; drug screening; gene therapy; antiinflammatory; antiasthmatic; antiarthritic; haemostatic; antiarteriosclerotic; immune disorder; cell culture; antiarthritic; haemostatic; antiarteriosclerotic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        haematopoiesis regulatory activity; tissue growth activity; immunomodulatory activity; activin- or inhibin-related activities; chemotactic or chemokinetic activities; haemostatic, thrombotic or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-457740/49
N-PSDB; ABA09151.
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27-APR-2000; 2000US-0560875.
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invention

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RESULT 14
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ID AAY48
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Best Local
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                                                                                                                                                                                                                                                                                        This invention decribes novel human nucleic acid sequences from tumor breast tissue which have cytostatic activity. The nucleic acid sequences can be used to produce and isolate full-length gene sequences. They can be used to express proteins, which can be used as tools to find an activity against breast cancer. The sequences can be used in sense or antisense form. They are especially useful for medicaments for gene therapy to treat breast cancer. Akv48540-y48617 represent protein fragments encoded by the expressed sequence tags described in the method
                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human nucleic acid sequences and tissue, useful for breast cancer
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treatment; tumour; cytostatic; medicament.
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                                                                                                                                                           Local
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                                                                                                                                                                                                                                                                        the invention.
                                                               MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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                                                                                                                                    Conservative
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pred. No. 2.8e-82;
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Similarity

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LALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKLGVKRCTDQMSL

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Best Local :
                                                                        The invention relates to a peptide comprising the amino acid sequences human lipophilin A, B, or C or its allelic variant or fragment comprisi at least one immunogenic epitope, which is purified and isolated, and m have the N-terminal acylated and/or C-terminal amidated or be a fusion protein. Molecules of the invention are used in the diagnosis and prognosis of mailgnant tumours associated with hormonally regulated organs such as uterus, ovary, prostate, testis, breast, kidney and thymus. This sequence represents the human Lipophilin B protein of the
                                                                                                                                                                                                                                          WPI; 200
N-PSDB;
                                                                                                                                                                                              Peptides having the sequence of human lipophilin A, ^{\circ} B and associated with carcinomas of hormonally regulated organs useful in the diagnosis and prognosis of various cancers
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                                           Sequence
                                                                                                                                                                          Claim 7; Fig 5; 22pp;
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           87.8%;
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"Encoded by ACG'
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2.2e-71;
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Perfect score:
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: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
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Copyright (c) 1993 - 2003 Compu
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) US-09-825-301-77

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US-09-985-911-4

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1 US-09-99-2598-260

1 US-09-99-335-260

9 US-09-99-344-260

9 US-09-99-436-260

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9 US-09-991-181-260

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Sequence 77, Appl
Sequence 29, Appl
Sequence 1, Appli
Sequence 4, Appli
Sequence 37, Appl
Sequence 27, Appl
Sequence 27, Appl
Sequence 260, App
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AL ESULT 1 S-09-825-301-77 Sequence 77, Application US/09825301 Patent No. US2002009738A1	58 12.9 162 58 12.9 226	5 13.8 178 5 13.2 95	64.5 14.3 100	9 75.5 16.8 93	91 20.2 105	6 111 24.7 74	5 142 31.6 . 74	3 172 38.2 69 4 171 38.0 112	2 172 38.2 69	1 202 44.9 90	0 202 44.9 90	9 206 45.8 111	7 238 52.9 83	6 238 52.9 83	5 238 52.9 83	4 238 52.9 83	3 238 52.9 83	2 238 52.9 83	1 238 52.9 83	0 238 52.9 83	
ALIGNMENTS	10 US-09-747-155-339 9 US-09-895-913A-362	l0 US-09-969-834-1 l0 US-09-925-299-1494		US-10-081-817-3	SOS	US-09-110-715-3	a (US-09-110-715-3	a	US-09-985-911-2	US-09-985-911-2	a c	US-09-990-455-	us-09-993-60-1-	SD	US-09-990-44:!-	US-09-991-07:I-	US-09-989-73::-	us-09-989-73::-26	LO US-09-989-72"-260	
	339, 7 362, Ap	Sequence 1, Appli Sequence 1494, Ap	Sequence 1430, Ap	32, A	Sequence 9, Appli Sequence 47, Appl	38,	35	8. %	12,	26,	25	sequence 5, Appli	260,	260,	260,	260,	•	260,	260,	Sequence 260, App	

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; LENGTH: 90
; TYPE: PRT
; ORGANISM: HOMO S
US-09-825-301-77
RESULT 2
US-09-110-716-29
; Sequence 29, Application US/09110716A
; Patent No. US20020034739A1
; GENERAL INFORMATION:
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APPLICANT: Zehentner, Barbara
APPLICANT: Zehentner, Barbara
APPLICANT: Persing, David H.
TITLE OF INVENTION: METHODS, COMPOSITIONS AND KITS FOR THE DETECTION
TITLE OF INVENTION: AND MONITORING OF BREAST CANCER
FILE REFERENCE: Z10121.513
CURRENT APPLICATION NUMBER: US/09/825,301
CURRENT FILING DATE: 2001-04-02
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 77
                                                                                                                                                                                                                                                                          Query Match 100.0%; Score 450; DB 10; Best Local Similarity 100.0%; Pred. No. 1.1e-45; Matches 90; Conservative 0; Mismatches 0;
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Molesh, David A.
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US-09-934-054-1
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CURRENT FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 41
SOFTWARE: PATENTIN VER: 2.0
SEQ ID NO 29
LENGTH: 90
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Best Local Similarity
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TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF CERTAIN TUMORS
FILE REFERENCE: 22000-20596.00
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                                                                                      INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS: LENGTH: 90 amino acids
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                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA
APPLICATION NUMBER: US/09/934,054
                                                                                                                                                                                         NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0077 US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                   FILING DATE: 21-Aug-2001 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hawkins, Phillip R.
TITLE OF INVENTION: BREAST TUMOR SPECIFIC PROTEINS
  SEQUENCE
                    MOLECULE
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                  TOPOLOGY: linear
ULE TYPE: <Unknow
                                                      TYPE: amino acid
STRANDEDNESS: double
                                                                                                                                                     TELEFAX: (650) 845-4166
                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/747,547 FILING DATE: <Unknown>
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Pred. No. 1.1e-45;
); Mismatches 0;
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US-09-985-911-4
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; ORGANISM: LpnB
US-09-110-716-37
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CURRENT FILING DATE: 2001-11-06
PRIOR APPLICATION NUMBER: 09/583,169
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR FILING DATE: 1999-03-08
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR FILING DATE: 1997-03-21
PRIOR FILING DATE: 1997-03-21
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Best Local S
Matches 90
                                     SOFTWARE: PatentIn Ver.
SEQ ID NO 37
LENGTH: 69
TYPE: PRT
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                                                                                                                                                                                                                                                                                                     Sequence 37, Application US/09110716A Patent No. US20020034739A1
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                                                                                                                 APPLICANT: Lehrer, Robert I.
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF CERTAIN TUMORS
FILE REFERENCE: 22000-20596,00
CURRENT APPLICATION NUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 41
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Best Local S
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: HUMAN ENDOMETRIAL SPECIFIC STEROID-BINDING FACTOR I, II AND II FILE REFERENCE: PF257D3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: human
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Length

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; LENGTH: 90
; TYPE: PRT
; ORGANISM: human
US-09-985-911-2
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US-09-985-911-2
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CURRENT FILING DATE: 2001-11-06
PRIOR APPLICATION NUMBER: 09/583,169
PRIOR FILING DATE: 2000-05-30
PRIOR APPLICATION NUMBER: 09/263,810
PRIOR FILING DATE: 1999-03-08
PRIOR FILING DATE: 1999-03-08
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR FILING DATE: 1997-03-21
                                             NUMBER OF SEQ ID NOS: 41
SOFTWARE: PatentIn Ver. 2
SEQ ID NO 27
LENGTH: 90
                                                                                                                                                                                                                                               Sequence 27, Application US/09110716A Patent No. US20020034739A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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APPLICANT: NI ET AL.
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Best Local :
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                                                                                                         APPLICANT: Lehrer, Robert I.
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC
FILE REFERENCE: 22000-20596.00
CURRENT APPLICATION NUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
ORGANISM: lipophilin A 09-110-716-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: HUMAN ENDOMETRIAL SPECIFIC STEROID-BINDING FACTOR I, II FILE REFERENCE: PF257D3
                                 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           82 VKILKKCSV 90
||||||||
61 VKILKKCSV 69
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                                                                                                                                                                                                                                                                                                                                                                  61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
les 53; Conserv
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Pred. No.
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Pred. No. 1.4e-25;
7; Mismatches 20;
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Best Local
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RESULT 8
US-09-864-761-44240
                                                                                                                                               NUMBER OF SEQ ID NOS: 49117
SOFTWARE: Annomax Sequence
SEQ ID NO 44240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/180,312 PRIOR FILING DATE: 2000-02-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Penn, Sharron G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT/US01/00667
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00664
PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: Aeomica-X-1
OTHER INFORMATION: MAP TO AC004127.1 OTHER INFORMATION: EXPRESSED IN LUNG, OTHER INFORMATION: EXPRESSED IN FETAL
                                                              ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             61 GVKRCTDQMSLQKRSLIAEVLVKILKKC
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                                                                                                                                                                                                                                                                                                 FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00670
FILING DATE: 2001-01-30
APPLICATION NUMBER: US 60/234,687
FILING DATE: 2000-09-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: PCT/US01/00668 FILING DATE: 2001-01-30 APPLICATION NUMBER: PCT/US01/00663
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FILLING DATE: 2000-09-27
APPLICATION NUMBER: PCT/US01/00666
FILING DATE: 2001-01-30
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APPLICATION NUMBER: GB
FILING DATE: 2000-10-04
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                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                     FILING DATE: 2000-06-30 APPLICATION NUMBER: US 09/774,203
                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/US01/00662 FILING DATE: 2001-01-30
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                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US 09/608,408
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: PCT/US01/00661
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Chen, Wensheng
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Pred. No. 1.(e-24;
2; Mismatches 23;
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SIGNAL - 1
LIVER, SIGNAL -
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OTHER INFORMATION: E
                                                                                                                       CURRENT APPLICATION NUMBER: 05/09/992,598
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/062787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/06250
PRIOR APPLICATION NUMBER: 60/065186
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
PRIOR FILING DATE: 1997-11-12
PRIOR PRIOR APPLICATION NUMBER: 60/065311
PRIOR APPLICATION NUMBER: 60/066770
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
PRIOR APPLICATION NUMBER: 60/075945
PRIOR APPLICATION NUMBER: 60/078910
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US-09-992-598-260
Sequence 260, Application US/09992598
Patent No. US20020160384A1
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TITLE OF INVENTION: Secreted
TITLE OF INVENTION: Acids Er
FILE REFERENCE: P2730P1C20
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         RR FILING DATE: 1998-04-28
RAPPLICATION NUMBER: 60/08
REFLING DATE: 1998-05-07
RR APPLICATION NUMBER: 60/08
RR APPLICATION NUMBER: 60/08
RR FILING DATE: 1998-05-28
RR FILING DATE: 1998-06-02
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Baker, Kevin P.
Botstein, David
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Tumas, D
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Gerritsen, Mary E.
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Stewart, Timothy A.
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Grimaldi, J.Christopher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Paoni, Nicholas F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Napier, Mary A.
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IN PLACENTA, SIGNAL = 1.2
IN BONE MARROW, SIGNAL = 1.1
IN ADULT LIVER, SIGNAL = 1.4
HIT: AW103293.1, EVALUE 5.00e-21
HIT: 095969, EVALUE 4.00e-22
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Pred. No. 2.9e-22;
); Mismatches 0;
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OR APPLICATION NUMBER: 60/088326
OR FILING DATE: 1998-06-04
OR APPLICATION NUMBER: 60/088167
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OR APPLICATION NUMBER: 60/088202
OR APPLICATION NUMBER: 60/088212
OR APPLICATION NUMBER: 60/088212
OR FILING DATE: 1998-06-05
OR APPLICATION NUMBER: 60/088217
OR APPLICATION NUMBER: 60/088217
OR APPLICATION NUMBER: 60/088217
OR APPLICATION NUMBER: 60/08825
OR APPLICATION NUMBER: 60/088655
OR FILING DATE: 1998-06-09
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DR FILLING DATE: 1998-06-10
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DR FILLWG DATE: 1998-06-23
DR APPLICATION NUMBER: 60/090429
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APPLICANT: Wood, William I.

APPLICANT: Zhang, Zemin
ITITLE OF INVENTION: Secreted and Transmemirane Pol
ITITLE OF INVENTION: Secreted and Transmemirane Pol
ITITLE OF INVENTION: Acids Encoding the Stime
FILE REFERENCE: P2730P1C6
CURRENT APPLICATION NUMBER: US/09/989,293%
CURRENT FILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: 60/049787
PRIOR APPLICATION NUMBER: 60/06250
PRIOR APPLICATION NUMBER: 60/06250
PRIOR APPLICATION NUMBER: 60/065186
PRIOR APPLICATION NUMBER: 60/065181
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OR APPLICATION NUMBER: 60/08/770
OR FILING DATE: 1997-11-24
OR APPLICATION NUMBER: 60/075945
OR FILING DATE: 1998-02-25
OR APPLICATION NUMBER: 60/078910
OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/083322
OR FILING DATE: 1998-04-28
OR APPLICATION NUMBER: 60/084600
OR FILING DATE: 1998-05-07
OR APPLICATION NUMBER: 60/0877106
OR FILING DATE: 1998-05-28
OR APPLICATION NUMBER: 60/087607
OR FILING DATE: 1998-05-07
OR FILING DATE: 1998-05-09
OR APPLICATION NUMBER: 60/087607
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/087609
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Godowski, Paul
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Roy, Margaret Ann
Stewart, Timothy
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Gerritsen, Mary E
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Pred. No. 4.4e-21;
3; Mismatches: 16
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DR FILING DATE: 1998-06-02

PARPLICATION NUMBER: 60/087827

DR FILING DATE: 1998-06-04

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PREPLICATION NUMBER: 60/088025

DR FILING DATE: 1998-06-04

PREPLICATION NUMBER: 60/088026

DR FILING DATE: 1998-06-04

PREPLICATION NUMBER: 60/088028

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PREPLICATION NUMBER: 60/088030

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PREPLICATION NUMBER: 60/088033

PREPLICATION NUMBER: 60/088036

PREPLICATION NUMBER: 60/088036

PREPLICATION NUMBER: 60/08826

PREPLICATION NUMBER: 60/08826

PREPLICATION NUMBER: 60/088212

PREPLICATION NUMBER: 60/088213

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R FILING DATE: 1998-06-19
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FILING DATE: 1998-07-01
APPLICATION NUMBER: 60/091519
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APPLICATION NUMBER: 60/091626
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     NUMBER: 60/092182
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NUMBER: 60/091633
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DR FILING DATE: 1998-06-17
DR APPLICATION NUMBER: 60/089600
DR FILING DATE: 1998-06-17
DR APPLICATION NUMBER: 60/089653
DR FILING DATE: 1998-06-17
DR APPLICATION NUMBER: 60/089801
DR FILING DATE: 1998-06-18
DR APPLICATION NUMBER: 60/089907
DR APPLICATION NUMBER: 60/089907
DR APPLICATION NUMBER: 60/089908

Query Match

Best Local Similarity

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Score Pred.

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Length 83;

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DATE: 1998-06-11

OR APPLICATION NUMBER: 60/088876

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OR APPLICATION NUMBER: 60/089105

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OR APPLICATION NUMBER: 60/089440

OR FILING DATE: 1998-06-16

OR APPLICATION NUMBER: 60/089512

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OR APPLICATION NUMBER: 60/089514

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OR APPLICATION NUMBER: 60/089599

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R FILING DATE: 1998-06-10
R APPLICATION NUMBER: 60/08
R FILING DATE: 1998-06-10
R APPLICATION NUMBER: 60/08
R ELING DATE: 1998-06-11
R APPLICATION NUMBER: 60/08

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APPLICATION NUMBER: 60/ FILING DATE: 1998-06-10

60/088824

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CURRENT FILING DATE: US/09/989,735
CURRENT FILING DATE: 2001-11-19
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-11-12
PRIOR FILING DATE: 1997-11-12
PRIOR FILING DATE: 1997-11-13
PRIOR FILING DATE: 1997-11-13
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PRIOR FILING DATE: 1997-11-3
PRIOR APPLICATION NUMBER: 60/066770
PRIOR APPLICATION NUMBER: 60/075945
PRIOR FILING DATE: 1998-02-25
PRIOR PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/083322
PRIOR APPLICATION NUMBER: 60/083322
PRIOR FILING DATE: 1998-04-28
PRIOR FILING DATE: 1998-04-28
PRIOR APPLICATION NUMBER: 60/084600
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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61 EVKHCTDQISFKKR 74
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                                                                                 APPLICATION NUMBER: 60/087607
FILING DATE: 1998-06-02
APPLICATION NUMBER: 60/087609
FILING DATE: 1998-06-02
  APPLICATION NUMBER: 60/087759
FILING DATE: 1998-06-02
APPLICATION NUMBER: 60/087827
FILING DATE: 1998-06-03
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Godowski, Paul J.
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Botstein, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wood, William I.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Napier, Mary A.
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Kljavin, Ivar J.
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OR FILING DATE: 1998-06-10
OR APPLICATION NUMBER: 60/088858
OR FILING DATE: 1998-06-11
OR APPLICATION NUMBER: 60/088861
OR FILING DATE: 1998-06-11
OR FILING DATE: 1998-06-11
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OR APPLICATION NUMBER: 60/088655
OR FILING DATE: 1998-06-09
OR APPLICATION NUMBER: 60/088734
OR FILING DATE: 1998-06-10
OR APPLICATION NUMBER: 60/088738
OR APPLICATION NUMBER: 60/088738
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60/091633

APPLICATION NUMBER: 60/090863 FILING DATE: 1998-06-26 APPLICATION NUMBER: 60/091360 FILING DATE: 1998-07-01

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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
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PRIOR APPLICATION NUMBER: 60/049787
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APPLICATION NUMBER: 60/078910
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Baker, Kevin P.
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Williams, P. Mickey
Wood, William I.
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Roy, Margaret Ann
Stewart, Timothy A.
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Gerber, Hanspeter
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OR APPLICATION NUMBER: 60/09

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APPLICATION NUMBER: 60/090472
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APPLICATION NUMBER: 60/091978
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Query Match 52.9%; Score 238; DB 9; Length 83; Best Local Similarity 60.8%; Pred. No. 4.4e-21; Matches 45; Conservative 13; Mismatches 16; Indels 0; Gaps

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RESULT 13
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APPLICANT: WOOD, William I.
APPLICANT: Zhang, Zemin
ITTLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C69
CURRENT PILING DATE: 2001-11-20
PRIOR APPLICATION NUMBER: 60/062250
PRIOR APPLICATION NUMBER: 60/062250
PRIOR APPLICATION NUMBER: 60/065186
PRIOR APPLICATION NUMBER: 60/065186
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Gerber, Hanspeter
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Botstein, David
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OR APPLICATION NUMBER: 60/08858
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DR APPLICATION NUMBER: 60/089512

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OR APPLICATION NUMBER: 60/089532

R APPLICATION NUMBER: 60/089532
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OR FILING DATE: 1998-06-04
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FILING DATE: 1998-06-25
APPLICATION NUMBER: 60/
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FILING DATE: 1998-06-24
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                                                                                                                                                                                                                                                        APPLICATION NUMBER: 60/091978
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FILING DATE: 1998-06-23
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                                                      1 Similarity 60.8 45; Conservative
                                                                                                                                                                                                                                          1998-07-07
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                                                                                                                     13;
                                                                                                                 Score 238; DB y; Pred. No. 4.4e-21;
                                                                                                                                              Length 83;
                                                                                                                    Indels
                                                                                                                    0,
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US-09-990-436-260
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/990,436 CURRENT FILING DATE: 2001-11-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/049787 PRIOR FILING DATE: 1997-06-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/PRIOR FILING DATE: 1997-10-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ashkenazi, Avi J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
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OR APPLICATION NUMBER: 60/07

OR FILING DATE: 1998-03-20

OR APPLICATION NUMBER: 60/08

OR FILING DATE: 1998-04-28

OR APPLICATION NUMBER: 60/08
                            R FILING DATE: 1998-06-04
R APPLICATION NUMBER: 60/0
R FILING DATE: 1998-06-04
R APPLICATION NUMBER: 60/0
R FILING DATE: 1998-06-04
                                                                                                                                                                                          R APPLICATION NUMBER: 60/01
R FILING DATE: 1998-06-03
R APPLICATION NUMBER: 60/01
R FILING DATE: 1998-06-04
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APPLICATION NUMBER: 60/
FILING DATE: 1998-05-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Watanabe, Colin K. Williams, P. Mickey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Grimaldi, J.Christopher Gurney, Austin L. Kljavin, Ivar J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wood, William I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Roy, Margaret Ann
Stewart, Timothy A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Paoni,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Napier, Mary A.
Pan, James
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gerritsen, Mary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fong, Sherman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eaton, Dan L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Desnoyers, Luc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Baker, Kevin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tumas,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Godowski,Paul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Goddard, Audrey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Application US/09990436
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                                                                                                                                                                                                                                                                                                                                                                          1998-06-02
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OR APPLICATION NUMBER: 05-04
(IOR FILING DATE: 1998-06-04
(IOR APPLICATION NUMBER: 60/08
TILING DATE: 1998-06-04
TILING TON NUMBER: 60/0
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NUMBER: 60/088030: 1998-06-04

OR APPLICATION NUMBER: 60/088326
DR FILING DATE: 1998-06-04
DR APPLICATION NUMBER: 60/088167
DR FILING DATE: 1998-06-05
DR APPLICATION NUMBER: 60/088202
DR FILING DATE: 1998-06-05
DR APPLICATION NUMBER: 60/088212
DR APPLICATION NUMBER: 60/088212
DR FILING DATE: 1998-06-05
DR FILING DATE: 1998-06-05

DR APPLICATION NUMBER: 60/088217

R FILING DATE: 1998-06-05

R APPLICATION NUMBER: 60/088655

R FILING DATE: 1998-06-09

DR APPLICATION NUMBER: 60/088734

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DR APPLICATION NUMBER: 60/088738

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R APPLICATION NUMBER: 60/090540

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APPLICATION NUMBER: 60/091519
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Pred. No. 4.4e-21;
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OR APPLICATION NUMBER: 60/088810
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US-09-991-181-260; Sequence 260, Applic; Publication No. US20; GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR FILING DATE: 1997-06-17
PRIOR FILING DATE: 1997-10-17
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PRIOR APPLICATION NUMBER: 60/065186
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PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/083322
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APPLICANT: Baker, Kevin P.
APPLICANT: Botstein, David
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APPLICATION NUMBER: 60/088167
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  Sequence 15, Application US/08912276
Patent No. 6183952
GENERAL INFORMATION:
APPLICANT: Billing-Medel, Patric
APPLICANT: Cohen, Maurice
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
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REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
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MEDIUM TYPE: 3.5 INCH DISKETTE
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE:
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ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER:
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OPERATING SYSTEM:
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TOPOLOGY: LI
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                                                                                                                                                                                                                                                                                   61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: amino acid
; STRANDEDNESS: sing
; TOPOLOGY: linear
; MOLECULE TYPE: No. 6
US-08-912-276-15
                                                                                                                                                                                                                               RESULT 4
US-09-583-169-4
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                                            GENERAL INFORMATION:
APPLICANT: Jian Ni, G
TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Loc
Matches
                                                                                                                                                                          Sequence 4, Application US/09583169 Patent No. 6338948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
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APPLICANT:
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NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,
                            CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 90 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for
CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Abbott Laboratories
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STREET: 100 Abbot:
CITY: Abbott Park
STATE: IL
                                                                                                                                                                                                                                                                                                                                61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                                                                                                                                                                                             61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                    MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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60064-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 90; DB 4; ilarity 100.0%; Pred. No. 3.4e-80; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roberts-Rapp, Lisa
Russell, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hodges, Steven C.
Klass, Michael R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Granados,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kratochvil, Jon D.
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CARELLA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                No. 6183952e
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                                                    Binding
27
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                                                                      Guo-Liang Yu and Reiner Gentz
Human Endometrial Specific St
Binding Factor I, II and III
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  BYRNE, BAIN, GILFILLAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US/08/912,276
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                                                                                                                                                                                                                                                                                                                                   90
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Steroid-

Length 90;

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Gaps

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CITY: ROSELAND STATE: NEW JERSEY

STREET:

ADDRESSEE: CECCHI, STEWART & OLSTEIN STREET: 6 BECKER FARM ROAD

COUNTRY:

USA

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; ORGANISM: Homo Sapiens US-09-215-818-6
                                                                                                                                     CURRENT FILING DATE: 1998-12-18
EARLIER APPLICATION NUMBER: 08/912,276
EARLIER FILING DATE: 1997-08-17
EARLIER APPLICATION NUMBER: 08/697,105
EARLIER FILING DATE: 1996-08-19
EARLIER APPLICATION NUMBER: 08/912,149
EARLIER FILING DATE: 1997-08-15
EARLIER APPLICATION NUMBER: 08/697,106
EARLIER APPLICATION NUMBER: 08/697,106
EARLIER FILING DATE: 1996-08-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-215-818-6
                                      SEQ ID NO 6
LENGTH: 90
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: COlpitts, Tracey
APPLICANT: COlpitts, Tracey
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
TITLE OF INVENTION: DETECTING DISEASES OF THE BREAST
FILE REFERENCE: 5972.US.P2
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 6, Application US/09215818A Patent No. 6379671
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                              CURRENT
                                                                                                  NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
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INFORMATION FOR SEQ ID NO: 4:
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LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 201-994-1700
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APPLICATION NUMBER:
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MEDIUM TYPE: 3.5 INCH DISKETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/215,818A FILING DATE: 1998-12-18
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                                                                                                  Windows Version 3.0
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ches 0;
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; STRANDEDNESS:
; TOPOLOGY: line
; MOLECULE TYPE: !
US-08-912-276-23
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US-08-912-276-23
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        Query Match 76.7
Best Local Similarity 100
Matches 69; Conservative
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Best Local :
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                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION: NAME: Becker, Cheryl L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
CITY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                               TYPE:
                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX:
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                                                                                                                                                                                                                                                                LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFF1.SEPLFKLSLAKFDAPPEAVAAKL 60
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60064-3500
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                                                                                                                                                                                                                                                              69 amino acids
                                                                                                                                                                                                                                                                                                                                                                                  847/938-2623
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Stroupe, Steven D.
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Klass, Michael R.
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                                                                                                                                                                                      linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM Compatible
                                                                                                                                                  No. 6183952e
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76.7%; 5cc.
100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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        s; Score 69; DB
s; Pred. NO. 6.2
0; Mismatches
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Pred. No.
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  DB 4; L. 6.2e-60; 0;
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                                                              Length 69;
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; MOLECULE TYPE: No. 6183952e
US-08-912-276-22
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Patent No.
                                                                                    Query Match
Best Local S
                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
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APPLICANT:
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APPLICANT:
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APPLICANT: Roberts-Rapp, Lisa
APPLICANT: Russell, John C.
APPLICANT: Stroupe, Steven D.
TITLE OF INVENTION: FOR DETECTING DISEASES OF THE BREAST
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: Fast-SEQ for Windows Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/912,276
                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 45 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                              NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 5972.US.P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
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MEDIUM TYPE: Diskette
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CITY: A
          61
                                                                                    Local Similarity
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ZIP: 60064-3500
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Abbott Park
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6183952
                                                                       Conservative
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Klass, Michael R.
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Colpitts, Tracey L.
Friedman, Paula N.
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                                                                                   50.0%;
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                                                                                   Score 45; Pred. No.
                                                                     Mismatches
                                                                                 DB 4; Lo
                                                                                                 Length 45
                                                                     Indels
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RESULT 9
US-08-912-276-16
; Sequence 16, Application US/08912276
; Patent NO. 6183952
; GENERAL INFORMATION:
; APPLICANT: Billing-Medel, Patric
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RESULT 8
US-08-912-276-21
                                                                                                                        ; TOPOLOGY: linear
; MOLECULE TYPE: No. 6183952e
US-08-912-276-21
                                                             Matches
                                                                                           Query Match
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                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
                                                                                                                                                                                                                                                              NAME: Becker, Cheryl L
REGISTRATION NUMBER: 5972
REFERENCE/DOCKET NUMBER: 5972
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
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APPLICANT:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: REAGENTS AND TITLE OF INVENTION: FOR DETECTING NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
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APPLICANT:
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SOFTWARE: FastSEQ for
CURRENT APPLICATION DATA:
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              69 MSLQKRSLIAEVLVKILKKCSV 90
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FILING DATE:
CLASSIFICATION:
                                                                                                                                                                    STRANDEDNESS:
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                                                            Local Similarity 100 les 22; Conservative
1 MSLQKRSLIAEVLVKILKKCSV 22
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No. 6183952
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Russell, John C.
Stroupe, Steven D.
VENTION: REACENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE BREAST
EQUENCES: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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Klass, Michael R.
Kratochvil, Jon D.
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Granados, Edward N.
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                                                                                                                                                                      single
                                                         24.4%; Score 22;
100.0%; Pred. No.
tive 0; Mismatc
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for Windows Version
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                                                             Mismatches
                                                                           DB 4; Le . 8.6e-15;
                                                                                        Length 22;
                                                             Indels
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Billing-Medel, Patricia A.

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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-912-276-16
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US-08-912-276-19
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                                                                                                                              Sequence 19, Application US/08912276
Patent No. 6183952
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                                                                                                                    GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 5977
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
TELEFAX: 847/938-2623
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SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NOTAGE
                       APPLICANT:
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APPLICANT:
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APPLICANT:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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                                                                                                      APPLICANT:
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                                                   APPLICANT:
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PPLICANT:
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ZIP: 60064-3500
                                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                              Similarity
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         Granados, Edward N. Hodges, Steven C. Klass, Michael R.
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Russell, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hodges, Steven C.
Klass, Michael R.
Kratochvil, Jon D.
Kratochvil, Jon D.
                                                   Gordon, Julian
                                                            Colpitts, Tracey L. Friedman, Paula N.
                                                                                         Cohen, Maurice
                                                                                                      Billing-Medel, Patricia
                                                                                                                                                                                                                                                                   Conservative
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Pred. No.
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US-08-912-276-17
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APPLICANT:
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           CORRESPONDENCE ADDRESS
                                    TITLE OF INVENTION:
                       NUMBER OF SEQUENCES:
                                                                                                                  APPLICANT:
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Roberts-Rapp, Lisa
Russell, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE BREAST

Abbott Laboratories

Hodges, Steven C. Klass, Michael R. Kratochvil, Jon D. Granados, Edward N.

Gordon, Julian Colpitts, Tracey L.

Friedman,

Paula N.

Billing-Medel, Patricia

0,

Maurice

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; TOPOLOGY: 11; MOLECULE TYPE: US-08-912-276-19
                                                                                       Query Match
                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
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Local
                                                                                                                                                                  LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 100 ._.
CTTY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM CO
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                            l Similarity
15; Conserv
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Russell, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASH'S OF THE BREAST
                                                            Conservative
                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IBM Compatible
                                                                                                                                       No. 6183952e
                                                                            100.0%;
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                                                                            Score 15;
Pred. No.
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nes 0;
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RESULT 12
US-08-912-276-18
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Best Local Similarity
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 5972
TELECOMMUNICATION INFORMATION:
TELEFAX: 847/935-1729
TELEFAX: 847/938-2623
TELEX:
VFORMATTON:
                                                                                                                              APPLICANT: Stroupe, S
TITLE OF INVENTION: F
TITLE OF INVENTION: F
NUMBER OF SEQUENCES:
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SEQUENCE CHARACTERISTICS
LENGTH: 16 amino acidd
            ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                     APPLICANT:
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                                                                                                                   CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                    APPLICANT:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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MEDIUM TYPE: Diskette
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  COMPUTER:
                                                  COUNTRY:
                                                                                                        ADDRESSEE:
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                                                                            r: 100 Abbott Park Road
Abbott Park
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RUSSEll, John C.
Stroupe, Steven D.
VENTION: REAGENTS AND METHODS USEFUL
VENTION: FOR DETECTING DISEASES OF THE BREAST
EQUENCES: 25
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Klass, Michael R.
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IBM Compatible
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100.0%; Pr
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Pred. No.
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RESULT 13
US-08-912-276-20
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; MOLECULE TYPE:
US-08-912-276-18
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Patent No. 6183952
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Best Local (
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                                                  SOFTWARE: FASTEM: DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICANT:
                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                     APPLICANT: Roberts-Rapp, Lisa
APPLICANT: Russell, John C.
APPLICANT: Stroupe, Steven D.
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL
TITLE OF INVENTION: FOR DETECTING DISEASES OF THE BREAST
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
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LENGTH: 16 amino acids
TYPE: amino acid
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REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
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                             CLASSIFICATION:
                                           FILING DATE:
                                                                                                                 COMPUTER:
                                                                                                                                                          ZIP: 60064-3500
                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                          STATE:
                                                                                                                                                                                                   CITY: Abbott Park
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SOFTWARE: FastSE(
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15; Conserv
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Klass, Michael R.
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Colpitts, Tracey L.
Friedman, Paula N.
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0; Mismatches
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4e-08;
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ATTORNEY/AGENT INFORMATION:

5972.US.P1

FILING DATE:

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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 16 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: NO. 6183952
US-08-912-276-20
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 Query Match
                                                                                                                                                                             APPLICATION UNMBER: US/08/821,451A
FILING DATE: MATCh 21, 1997
CLASSIFICATION : 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/014,724
APPLICATION NUMBER: 60/014,724
FILING DATE: MATCh 21, 1996
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J. 6.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
TELEPAN: 201-994-1744
TURNOBLATION FOR SECT IT NO. 2.
                                                                                                                              TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Jian N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCI
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Becker, Cheryl L
REGISTARION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/935-1729
                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific Steroid-
TITLE OF INVENTION: Binding Factor I, II and III
                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        69 MSLQKRSLIAEVLVK 83
                                                                                         STRANDEDNESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: UZIP: 07068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: ROSELAND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, ADDRESSEE: CECCHI, STEWART & OLSTEIN STREET: 6 BECKER FARM ROAD
                                                                            TOPOLOGY:
                                                                                                             TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 847/938-2623
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E: No. 6183952e
                                                                            LINEAR
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100.0%;
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 Score 8;
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US-09-263-810-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLITAMENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 32,073
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MEDIUM TYPE: 3.5 INCH DISKETTE
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LENGTH: 90 AMINO ACIDS
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APPLICATION NUMBER:
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CLASSIFICATION:
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STATE: NEW JERSEY
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	2000 (Rel. 39, Crea 2000 (Rel. 39, Last	
E 15	(Rel. 41, Last annotation upda	
	Chordata;	
X NC	<pre>I_TaxID=9606;</pre>	
	[1] SEQUENCE FROM N.A.	
	MEDLINE-99167354; PubMed-10066439;	
	"Lipophilins: human peptides homologous to rat prostatein.";	
	1- FUNCTION: MAY BIND ANDROGENS AND OTHER STEROIDS, MAY ALSO BIND	
ឥឥ	ESTRAMUSTINE, A CHEMOTHERAPEUTIC AGE: T USED FOR PROSTATE CANCER. MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES.	
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	This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -	
	opean Bioinformatics Institute. There are no res	

Ϋ́ Query Match Best Local S Matches 90 Interpro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
SMART; SM00096; UTG; 1.
Signal.
SIGNAL 21 POTENT CHAIN 22 90 LIPOPH SEQUENCE 90 AA; 9925 MW; 17BB55 the European Bioinformatics Institute. There are no restrictions on its monomerce by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (see http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch). EMBL; AJ224172; CAA11864.1; -. Genew; HGNC:18396; SCGB1D2. l Similarity 90; Conserv Conservative 100.0%; 0; POTENTIAL. LIPOPHILIN B. 17BB555ED0 \SD1AF CRC64; Score 90; DB 1; I Pred. No. 3.1e-84; ; Mismatches 0; Length 90 Indels 0, Gaps 60 0

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MEDLINE-93009001; PubMed-1395029;
MEDLINE-93009001; PubMed-1395029;
Bernard A., Roels H., Lauwerys R., Witters R., Gielens C.,
Soumillion A., Van Damme J., De Ley M.;
Soumillion A., Van Damme J., De Ley M.;
"Human urinary protein 1: evidence for identity with the Clara
"Human urinary protein 1: respiratory tract and urogenital
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01-0CT-1989 (Rel. 12, Created)
01-0CT-1989 (Rel. 12, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Clara cell phospholipid-binding protein precursor (CCPBP) (Clara cell phospholipid-binding protein precursor (CCPBP) (Clara cell phospholipid-binding protein (Urine protein I)
                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-93016476; PubMed-1400743; Okutani R., Itoh Y., Hirata H., Kasahara T., Mukaida N., Kawai T.; "Simple and high-yield purification of urine protein 1 using immunoaffinity chromatography: evidence for the identity of urine protein 1 and human Clara cell 10-kilodalton protein.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Beato M., Suske G.; "Human CC10, the homologue of rabbit uteroglobin: genomic cloning, chromosomal localization and expression in endometrial cell lines."; Hum. Mol. Genet. 1:371-378(1992).
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MEDLIND-89000784; PubMed-3167058;
Singh G., Katyal S.L., Brown W.E., Phillips S., Kennedy A.L.,
Anthony J., Squeglia N.;
*Amino-acid and cDNA nucleotide sequences of human Clara cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE OF 1-18 FROM N.A. MEDLINE-93250776; PubMed-1284526; Wolf M., Klug J., Hackenberg R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hay J.G., Danel C., Chu C., Crystal R.G.;
"Human CC10 gene expression in airway epithelium and subchromosomal locus suggest linkage to airway disease.";
Am. J. Physiol. 268:L565-L575(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE-95250987; PubMed-7733299; Hay J.G., Danel C., Chu C., Cryst
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EMBL; U01101; AAA81885.1; -.
EMBL; U01102; AAA18297.1; -.
EMBL; BC004481; AAH04481.1; -.
EMBL; X59875; CAA42532.1; -.
EMBL; X59875; CAA42532.1; -.
FIR; JS0036; JS0036.
PIR; S26651; S26651
PIR; S26651; S26651
PIR; S26651; S26651
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InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "Structure of a human Clara cell phospholipid-binding protein-ligand complex at 1.9-A resolution.";
Nat. Struct. Biol. 1:538-545(1994).
-I- FUNCTION: BINDS PHOSPHATIDYLCHOLINE, PHOSPHATIDYLINOSITOL, POLYCHLORINATED BIPHENYLS (PCB) AND WEAKLY PROGESTERONE, POTENT INHIBITOR OF PHOSPHOLIPASE A2.
-I- SUBUNIT: Homodimer; antiparallel disulfide-linked.
-I- SUBUNIT: Homodimer; antiparallel disulfide-linked.
-I- SUBUNIT: BELONGS TO THE ULLMONARY AIRWAYS).
                                                                                                                                                                                        SEQUENCE
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                        PSC2_RAT
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Proteomics 2:112-120(2002).
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7 -> A (IN DBSNP:1802632).
/FTId=VAR_012046.
C -> E (IN REF. 7).
FE65ACA678F12ABD CRC64;
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Mammalia; Eutheria;
NCBI_TaxID=10116;
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peptide (2).
Rattus norvegicus (Rat).
Rattus norvegicus (Chordata;
                                                                                                                                                                                                   EMBL; X05034; CAA28708.1; -.
EMBL; V01256; CAA24569.1; -.
EMBL; J00776; AAA51641.1; -.
PIR; A03251; BORT2.
PIR; A26671; A26671.
                                                                                                                                                                                                                                                                                              the European Bioinformatics Institute. There are no restrictions use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for commodified and this statement is not removed.
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                                                                                                                                           Steroid-binding; SIGNAL 1
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MEDLINE-83209619; F
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                                                          CONFLICT
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"Prostatic steroid binding prote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Parker M., Needham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-82220075; PubMed-6896362;
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                                                                                                                                                                                                                                                                                                                                                                              GLYCOPROTEIN OF VENTRAL PROSTATE GLAND.

GLYCOPROTEIN OF VENTRAL PROSTATE GLAND.

GUBUNIT: PROSTATEIN IS COMPOSED OF THREE DIFFERENT PEPTIDES C1, C2 AND C3:C3 (S) HETERODIMERS WHOSE NONCOVALENT ASSOCIATION FORMS TETRAMERIC (C1:C3/C3:C2) PROSTATEIN MOLECULES.

GUBCELLULAR LOCATION: Secreted.

PTM: LINKED BY THREE DISULFIDE BONDS TO C3.

MISCELLANEOUS: THE HETERODIMER CAN BIND NON-POLAR STEROIDS, CHOLESTEROL AND A GROUP OF SMALL PROLINE-RICH PEPTIDES.

SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FUNCTION: PART
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Acids Res. 15:1627-1641(1987).
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L., Decourt J.-L.,
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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01-AUG-1988 (Rel.
16-OCT-2001 (Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence analysis.";
Biochemistry 27:4214-4221(1988).
-i-FUNCTION: SPERMINE-BINDING PROTEIN IS AN
VENTRAL PROSTATE GLYCOPROTEIN THAT BINDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-87137538; PubMed-3818623; Chang C., Saltzman A.G., Hiipakka
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Pfam; PF01419; Jacalin;
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Anderegg R.J., Carr S.A., Huang I.-Y.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE, AND REVISIONS
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LLLVTLAL
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                                                                                                              8; Conserv
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                                                                                                                 Score 8; D3 1
Pred. No. 1.2
0; Mismatches
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ASP/GLU-RIC1 (ACIDIC).
3BEB01A02.4517A65 CRC64;
                                                                                                                                                                                                                                                                                                                                                            PROSTATIC SPERMINE-BINDING
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Sciurognathl; Muridae;
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of tandem
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and nucleotide sequence
control of mRNA level.";
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              HBA_BALAC STANDARD; PRT; 141 AA P18971; 01-NOV-1990 (Rel. 16, Created) 01-NOV-1990 (Rel. 16, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hemoglobin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Lipophilin, a novel heterodimeric protein of human tears.";
FEBS Lett. 432:163-167(1998).
-!- FUNCTION: MAY BIND ANDROGENS AND OTHER STEROIDS, MAY ALSO BIND
ESTRAMUSTINE, A CHEMOTHERAPEUTIC AGENT USED FOR PROSTATE CANCER.
MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES.
-!- SUBUNIT: HETERODIMER OF A LIPOPHILIN A AND A LIPOPHILIN C
(MAMMAGLOBIN B) MONOMER ASSOCIATED HEAD TO HEADD.
-!- SUBCELLULAR LOCATION: Secreted (Potential).
-!- TISSUE SPECIFICITY: EXPRESSED IN LACHRYMAL GLAND, THYMUS, KIDNEY
TESTIS, OVARY AND SALIVARY GLAND.
-!- SUBCELLULAR LOCATION: SECRETARY GLAND.
-!- SIBLARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Lipophilin A precursor (Secretoglobin family 1D member SCGBIDI OR LIPHA.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Lehrer R.I., Xu G.,
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PF01099; Uteroglobin; 1.
S; PR00486; UTEROGLOBIN.
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                   annotation update)
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Pred. No.
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DF2D4F7565A87D34 CRC64;
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RESULT 7
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P01963;
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SEQUENCE
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Cetacea) hemoglo
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Balaenoptera acutorostrata (Minke whale) (Lesser rorqual, Balaenoptera ecutorostrata; Craniata; Vertebrata; Eute.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute.
Mammalia; Eutheria; CetartLodactyla; Cetacea; Mysticeti;
                   Pfam; PF00042; globin; 1.
PRINTS; PR00612; ALPHAHAEM.
                                                                                                                                                                     -i- FUNCTION: Involved in oxygen transport from various peripheral tissues.
-i- SUBUNIT: Heterotetramer of two alpha chains
                                                                                                                                                                                                                   (Ceratotherium simum, perissodactyla): beta 2 Glu., Hoppe-Seyler's Z. Physiol. Chem. 363:1077-1085(192)-1- FUNCTION: Involved in oxygen transport from the
                                                                                                                                                                                                                                                                   MEDLINE-83055102; PubMed-7141412;
Mazur G., Braunitzer G., Wright P.G.;
"The primary structure of the hemoglobin
                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   HBA.
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21-JUL-1986 (Rel. 01, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation updat
Hemoglobin alpha chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- TISSUE SPECIFICITY: RED BLOOD CELLS.
-!- SIMILARITY: BELONGS TO THE GLOBIN FA
PIR; S06520; HAWHK.
HSSP; P01922; 1BZ0.
InterPro; IPR002338; Alpha_haem.
InterPro; IPR000971; Globin.
                                                         InterPro; IPR002338; Alpha_haem
InterPro; IPR000971; Globin.
                                                                                                                                   -!- TISSUE SPECIFICITY:
                                                                                                                                                                                                                                                                                                                                     SEQUENCE
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                                                                                               A02286; HARNW
; P01958; 2MHB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oxygen transport; Transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58
87
21
96
141 /
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
GLOBIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                     n (White rhinocero;
; Chordata; Crania;
; Perissodactyla;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   58
87
21
96
15406
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7.8%;
                                                                                                                                 Red blood cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W.
                                                                                                                                                                                                                                                                                                                                                                                                         rhinoceros) (Square-lipped rhinoceros).
ta; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
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Pred. No
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IRON
G ->
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0B0C26CDF7B72B53 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Erythrocyte;
                                                                                                                                                                                                                                                                                                                                                                                       Rhinocerotidae; Ceratotherium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Zaidi Z.H., Braunitzer
e (Balenoptera acutorost
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DISTAL LIGAND).
PROXIMAL LIGAND).
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                                                                                                                                                                                                                                                     Glu.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 141
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                                                                                                                                                                                                             lung
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Euteleostomi;
                                                                                                                                                                                                                                                                         rhinoceros
                                                                                                                                                                           beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            beta chains.
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                                                                                                                                                                           chains
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HBA_RHIUN
ID HBA_R
P0990
AC P0990
DT 01-MA
DT 15-JU
DE Hemog
GN HBA.
OS Rhino
OC Eukar
OC Mammaa
OX NCBI_
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Best Local
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                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P01949;
21-JUL-1986 (Rel. 0
21-JUL-1986 (Rel. 0
15-JUN-2002 (Rel. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ERIEU
                                                             P09906;
01-MAR-1989 (Rel. 10, Created)
01-MAR-1989 (Rel. 10, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                            METAL
SEQUENCE
          Rhinoceros unicornis (Greater Indian rhinoceros).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Perissodactyla; Rhinocerotidae;
                                                                                                                                                                                                                                                                                       PRINTS; PRO0612; ALPHAHAEM.

PROSITE; PS01033; GLOBIN; 1.

Heme; Oxygen transport; Transport; Erythrocyte.
                                                                                                                                                                                                                                                                                                                                                               PIR; A02271; HAHH.
HSSP; P01922; 1BZ0.
                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=79109529; PubMed=762046;
Maita T., Araya A., Matsuda G., Goodman M.;
"Amino acid sequences of the alpha and beta cha
hemoglobin of the European hedgehog, Erinaceus
J. Biochem. 85:259-269(1979)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Erinaceus europaeus (Western European hedgehog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Insectivora; Erinaceidae; Erinaceinae;
                                                       Hemoglobin alpha
                                                                                                                                                                                                                                                                                                                             Pfam; PF00042; globin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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                                                                                                              HBA_RHIUN
                                                                                                                                                                                                                                                                                                                                        InterPro; IPR002338; Alpha_haem.
InterPro; IPR000971; Globin.
                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- FUNCTION: Involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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                                                                                                                                                                      105
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105 LLVTLAL 111
                                                                                                                                                                                                                           Local
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                                                                                                                                                                                          8 LLVTLAL 14
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                                                                                                                                                                      LLVTLAL 111
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                                                                                                                                                                                                                Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 7; Conserv
                                                                                                                                                                                                                                                           58
87
141 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      141 AA;
                                                                                                                                                                                                              7.8%; ilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transport;
                                                                                                              STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
                                                        chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01, Last sequence update)
41, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01, Created)
                                                                                                                                                                                                                                                            15122 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                in oxygen transport from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Transport; Erythrocyte.
IRON (HEME DISTAL LIGAND).
IRON (HEME PROXIMAL LIGAND).
                                                                                                                                                                                                              Score 7; DB 1; Pred. No. 7.1
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 7; I
                                                                                                                                                                                                                                                           IRON (HEME DISTAL LIGAND).
IRON (HEME PROXIMAL LIGAND).;
E4EE4DE6485050F6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Goodman M.;

pha and beta chains of adult

----- ouropaeus.";
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                                                                                                              141
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                                                                                                                                                                                                                        DB 1;
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. 7.1;
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                                                                                                                                                                                                                                   Length 141;
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                                                                                                                                                                                                                                                                                                                                                                                                            and
                                                                                                                                                                                                                                                                                                                                                                                                                                the
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                                                                                                                                                                                                                Indels
          Euteleostomi;
Rhinoceros.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Euteleostomi;
ceinae; Erinaceus.
                                                                                                                                                                                                                                                                                                                                                                                                           two beta chains
                                                                                                                                                                                                                                                                                                                                                                                                                                lung to the
                                                                                                                                                                                                               0
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HBA_TAFTE
ID HBA_T
AC POL196
AC POL1
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       P01962;
21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence upda:e)
21-JUL-1986 (Rel. 01, Last sequence upda:e)
15-JUN-2002 (Rel. 41, Last annotation up Mate)
15-JUN-2002 (Rel. 01, Last sequence upda:e)
15-JUN-2002 
                                                                                              InterPro; IPR002338; Alpha_haem.
InterPro; IPR000971; Globin.
Pfam; PF00042; globin; 1.
PRINTS; PR00612; ALPHAHAEM.
PROSITE; PS01033; GLOBIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the beta chains.";
Hoppe-Seyler's Z. Physiol. Chem.
-i-FUNCTION: Involved in oxygen
various peripheral tissues.
-i-SUBUNIT: Heterotetramer of tv
-i-TISSUE SPECIFICITY: Red blood
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Heme; Oxygen transport; Transport;
METAL 58 58 IRON (
METAL 87 87 IRON (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pfam; PF00042; globin; 1.
PRINTS; PR00612; ALPHAHAEM.
PROSITE; PS01033; GLOBIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Abbasi A., Weber R.E., Braunitzer G., Goltenboth R.; "Molecular basis for ATP/2,3-bisphosphoglycerate control switch-over (poikilotherm/homeotherm) an intermediate amino-acid sequence in the hemoglobin of the great Indian rhinoceros (Rhinoceros unicornis,
                                      Polymorphism
                                                                  Heme; Oxygen transport;
                                                                                                                                                                                                                                                                      PIR; A02285; HATPI.
PIR; A91727; HATP2B.
HSSP; P01958; 2MHB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Perissodactyla: the primary structure cf hemoglobins from the lowland tapir (Tapirus terrestris): glutamic acid in position 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mazur G., Braunitzer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=85053021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HBA_TAPTE
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HSSP; P01958; 1IBE.
InterPro; IPR002338; Alpha_haem.
InterPro; IPR000971; Globin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Perissodactyla).";
Biol. Chem. Hoppe-Seyler 368:323-332(1987)
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SEQUENCE.
MEDLINE=87271129; PubMed=3606819;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 LLVTLAL 14
                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: BELONGS TO
                                                                                                                                                                                                                                                                                                                                                                                                                                          POLYMORPHISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: Involved in oxygen transport from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LLVTLAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               141 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PubMed-6149994;
      58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15319 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          7.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        in oxygen transport from
                                                                     Transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ARE
                                                                                                                                                                                                                                                                                                                                                                           THE GLOBIN FIMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                       of two alpha chains and two beta chains blood cells.
TWO ALLELES. THE SEQUENCE SHOWN IS THAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 7; I
IRON (HEME DISTAL LIGAND).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ort; Erythrocyte.

RON (HEME DISTAL LIGAND).

RON (HEME FROXIMAL LIGAND).

9612DA629640B260 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        365:1097-1106(1984).
                                                                  Erythrocyte;
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P54158;
01-OCT-1996
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VARIANT
SEQUENCE
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HSSP; P01922; 1BZO.
InterPro; IPR002338; Alpha_haem.
InterPro; IPR000971; Globin.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti; Delphinidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1990 (Rel. 16, Created)
01-NOV-1990 (Rel. 16, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
                                                                                                                                                         Hypothetical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRINTS; PRO0612; ALPHAHAEM. PROSITE; PS01033; GLOBIN; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Tursiops truncatus, Cetacea).";
Biomed. Biochim. Acta 42:685-695(1983).
-!- FUNCTION: Involved in oxygen transport from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P18978;
01-NOV-1990
                                                       Bacteria;
                                                                                        Bacillus subtilis
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7; Conserv
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141 AA;
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D -> G (IN ALPHA-2).
; E3727EB6650C5BD1 CRC64;
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(HEME PROXIMAL LIGAND).
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RA Kunst F. (Ogasawara N., Moszer I., Albertini A.M., Alloni G.,
RA Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert S.,
RA Borniss R., Boursier L., Brans A., Barun M., Brignell S.C., Bron S.,
RA Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,
RA Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Choi S.K., Edwin J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
RA Chim K.D., Errington J., Fabret C., Ferrari E., Foulger D.,
RA Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N.,
RA Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N.,
RA Guiseppi G., Guy B.J., Haga K., Halcch J., Harwood C.R., Henaut A.,
RA Hilbert H., Holsappel S., Hosono S., Hullo M.F., Itaya M., Jones L.,
RA Guiseppi G., Guy B.J., Haga K., Halcch J., Harwood C.R., Henaut A.,
RA Kobayashi Y., Koetter P., Koningstein G., Krogh S., Kumano M.,
RA Kobayashi Y., Koetter P., Koningstein G., Krogh S., Kumano M.,
RA Medina N., Levine A., Liu H., Masuda S., Mauel C., Medigue C.,
RA Lee S.M., Levine A., Liu H., Masuda S., Mauel C., Medigue C.,
RA Presecan E., Pujic P., Purnelle B., Ropen M., Sadaie Y.,
RA Parro V., Pohl T.M., Portetelle D., Porrwollik S., Prescott A.M.,
RA Parro Y., Pujic P., Purnelle B., Roche B., Rose M., Sadaie Y.,
RA Rieger M., Rivolta C., Rocha E., Roche B., Rose M., Sadaie Y.,
RA Rauchi J., Sekowska A., Seror S.J., Serror P., Shin B.S., Soldo B.,
RA Sato T., Scandan E., Schleich S., Schroeter R., Scoffone F.,
RA Satoria A., Tanakoshi A., Tanaka T., Takahashi H., Takemaru K.,
RA Yari A., Wambutt R., Wedler E., Wedler H., Weitzenegger T.,
RA Yari A., Wambutt R., Wedler E., Wedler F., Vasumoto K., Yata K.,
RA Yoshida K., Yoshikawa H.F., Zumsten E., Yoshikawa H., Danchin A.,
RThe complete genome sequence of the Gram-positive bacterium Bacillus
                                    Hypothetical
TRANSMEM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
STRAIN-168 / Marburg;
MEDLINE-96349105; PubMed-8760912;
MEDLINE-96349105, PubMed-8760912;
                                                                                                                                                           EMBL;
EMBL;
                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chen L., James L.P., Helmann J.D.;
"Metalloregulation in Bacillus subtilis: isolation and characterization of two genes differentially repressed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 155-168 FROM N.A. MEDLINE=93374837; Pubmed=8396117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Sequence analysis of the Bacillus subtilis chromosome region between the serA and kdg loci cloned in a yeast artificial chromosome."; Microbiology 142:2005-2016(1996).
                                                                                                                      SubtiList;
                                                                                                                                         EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Serror P.;
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-!- SUBCELLULAR LOCATION: Integral membrane protein (Potential).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-98044033; PubMed-9384377;
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Z99115; CAB14121.1; -.
L19548; -; NOT_ANNOTATED_CDS.
Z22929; -; NOT_ANNOTATED_CDS.
                                                                                                                      BG11603;
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               protein;
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n; Transmembrane; Co.
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30-MAY-2000
30-MAY-2000
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SEQUENCE FROM N.A.
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MEDLINE-97061201; Pubmed-8905231;
Kaneko T., Sato S., Kotani H., Tanaka A., Asamizu E., Nakamura Y.,
Kaneko T., Sato S., Kotani H., Tanaka A., Asamoto S., Kimura T.,
Miyajima N., Hirosawa M., Sugiura M., Sasamoto S., Kimura T.,
Miyajima N., Matsuno A., Muraki A., Nakazaki N., Naruo K., Okumura
Hosouchi T., Matsuno A., Muraki A., Nakazaki N., Yamada M., Yasuda M.,
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15-JUL-1998 (Rel. 36, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Sequence analysis of the genome of the unicellular cyanobacterium synechocystis sp. strain PCC6803. II. Sequence determination of the entire genome and assignment of potential protein-coding regions."

DNA Res. 3:109-136(1996).
                                                                                                                                                                                                                                                                                                                                                                                    METTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synechocystis sp. (strain PCC 6803).
Bacteria; Cyanobacteria; Chroococcales; Synechocystis.
                       SEQUENCE FROM N.A. STRAIN-Delta H;
                                                                                                                                               Methanobacterium thermoautotrophicum.
Archaea; Euryarchaeota; Methanobacteria;
                                                                                                                                                                                                   PCRB protein homolog. PCRB OR MTH552.
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MEDLINE=98037514; PubMed=9371463;
                                                                                                NCBI_TaxID=187420;
                                                                                                                          Methanobacteriaceae;
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(Rel. 39, Last sequence update)
(Rel. 40, Last annotation updat
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0457 MW; F4B40D995AFD2138 CRC64;
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STRAIN=KIZ MG1655;
MEDLINE=97426617; PubMed=9278503;
MEDLINE=97426617; PubMett G. III, Bloch C.A., Perna N.T., Burland Blattner F.R., Plunkett G. III, Bloch C.A., Rode C.K., Mayhew G.F. Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew G.F. Riley M., Collado-Vides J., Glasner J.D., Rode M.A., Rose D.J., T. Davis N.W., Kirkpatrick H.A., Goeden M.A., Rose D.J.,
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                         Itoh T., Aiba H., Baba T., Fujita K., Hayashi K., Inada T., Kasai H., Kimura S., Kitakawa M., Kitagana M., Makino K., Mi Mizobuchi K., Mori T., Motomura K., Nakade S., Nakada S., Sel Sivasundaram S., Tayami H., Takeda J., Takemoto K., Wada C., Yamamoto Y., Horiuchi T., Yanamoto Y., Horiuchi T., Takeda J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Pakada J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Pakada J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Pakada J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Pakada J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Pakada J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Pakada J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Pakada J., Takemoto K., Wada C., Yanamoto Y., Horiuchi T., Yanamoto Y., Hor
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InterPro; IPR003009; FMN_enzyn
InterPro; IPR002911; PcrB.
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Putative 1-aminocyclopropane-1-carboxylate deaminase
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15-JUL-1998 (Rel. 36, Created)
15-JUL-1999 (Rel. 38, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                             MEDLINE=97251358; PubMed=9097040;
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                                                                                                                                                                                                                                                                                                                              complete genome sequence nce 277:1453-1474(1997).
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sequence of the Escherichia to the 40.1-50.0 min region
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on the linkage map.";
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                                                                            ei G., Seki :, Wada C.,
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Miki T.,
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Job time : 14 secs
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Best Local Similarity 100.0%; P
Matches 7; Conservative 0;
                                                                                                                                                                                                                          EMBL; AE000284; AAC74986.1; ALT_INIT.
EWBL; D90832; BAA15746.1; ALT_INIT.
EMBL; D90833; BAA15746.1; ALT_INIT.
ECOGene; EG14038; yedo.
InterPro; IPR001926; B6_enzyme_beta.
Pfam; pF00291; PALP; 1.
TIGRFAMs; TIGR01275; ACC_deam_rel; 1.
Hypothetical protein; Lyase; Pyridoxal phosphate; Complete proteome.
Hypothetical protein; Lyase; Pyridoxal phosphate (BY SIMILARITY).
SEQUENCE 328 AA; 35153 MW; 4179DE645C0B32D8 CRC64;
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-I- COFACTOR: PYRIDOXAL PHOSPHATE (BY SIMILARITY).
-I- SIMILARITY: BELONGS TO THE ACC DEAMINASE FAMILY.
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-1- CATALTIC ACTIVITY: 1-aminocyclopropane-1-carboxylate + H(2)0 = 2-
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Delius H., Darai G., Fluegel R.M.;

Delius H., Darai G., Fluegel R.M.;

"DNA analysis of insect iridescent virus

permutation and terminal redundancy.";

J. Virol. 49:609-614(1984).
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Fischer M.; Schnitzler P., Delius H., Darai G.;

"Identification and characterization of the repetitive the genome of insect iridescent virus type 6.";
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Virology 160:66-74(1987).
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Viruses; dsDNA viruses, no RNA stage; Iridoviridae; Iridovirus
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MEDLINE=87321126; PubMed=2820141;

Schnitzler P., Soltau J.B., Fischer M., Reisner H., Scholz
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Lorbacher de Ruiz H., Gelderblom H., Hofmann V
"Insect iridescent virus type 6 induced toxic
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DEPTH OF THE SEQUENCE OF CALLO INTESSECRET VITUS BETWEEN COORDINATES 0.101 and 0.391; similarities in coding insect and vertebrate indoviruses.";

Virus Genes 15:235-245(1997).
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SEQUENCE FROM N.A.
MEDLINE-94292906; PubMed-8021587;
Sonntag K.C., Muller
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SEQUENCE FROM N.A.
MEDLINE=94167241; PubMed=8121799;
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Stohwasser R., Raab K., Schnitzler P., Janssen W., Darai G.;
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*Insect iridescent virus type 6 encodes a pol
largest subunit of eukaryotic RNA polymerase
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Delius H., Darai C.;
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                       iridescent virus.";
Genes 17:243-258(1998).
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Zhao C., Nguyen T.X., Lehrer R.I.;
Zhao C., Nguyen T.X., Lehrer R.I.;
"Rabbit Lipophilins.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ da:
EMBL; AF308616; AAG42804.1;
Interpro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
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                      Cepaea nemoralis Mitochondrion.
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Jakob N.J., Muller K., Bahr U., Darai G.;
"Analysis of the First Complete DNA Sequence
Iridovirus: Coding Strategy of the Genome of
Virology 286:182-196(2001).
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8; Conser
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8; Conser
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Last sequence up
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Pred. No.
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                                              snail)
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Iridescent Virus.";
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Mollusca; Gastropoda; Pulmonata;

Stylommatophora;

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RESULT
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Q92QJ8;
Q1-DEC-2001 (TrEMBLrel. 19, Created)
Q1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
Q1-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Q1-MAR-2002 (TREMBLREL 20, Last annotation update)
                                                                MEDLINE-21396507; PubMed=11481430;
Capela D., Barloy-Hubler F., Gonzy J., Bothe G., Ampe F., Batut J., Gapela D., Barloy-Hubler F., Gonzy J., Cadieu E., Dreano S., Gloux S., Boistard P., Becker A., Boutry M., Cadieu E., Dreano S., Gloux S., Godrie T., Goffeau A., Kahn D., Kiss E., Lelaure V., Masuy D., Pohl T., Portetelle D., Puehler A., Purnelle B., Ramsperger U., Renard C., Thebault P., Vandenbol M., Weidner S., Galibert F., "Analysis of the chromosome sequence of the legume symbiont Sinorhizobium meliloti strain 1021.";
Proc. Natl. Acad. Sci. U.S.A. 98:9877-9882(2001).
EMBL, AL591786; CAC45901.1; -.
InterPro; IPR000063; Thiored.
PROSITE; PS00194; THIOREDOXIN; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Evolution of pulmonate gastropod mitochondrial genomes: compari of complete gene organizations of Euhadra, Cepaea and Albinaria implications of unusual tRNA secondary structures."; Genetics 145:749-758(1997).

EMBL; U23045; AAC09520.1;
EMBL; U23045; AAC09520.1;
InterPro; IPR001505; Copper_CuA.
InterPro; IPR002088; PPTA.
InterPro; IPR002088; PPTA.
InterPro; IPR000822; Znf_C2H2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rhizobium meliloti (Sinorhizobium meliloti). Bacteria; Proteobacteria; alpha subdivision; Rhizobiaceae; Sinorhizobium.
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Pfam; PF02790; COX2_TM; 1.
Pr0Dom; PD000131; Copper_CuA; 1.
PROSITE; PS00904; PPTA; UNKNOWN_1.
PROSITE; PS00028; ZINC_FINGER_C2H2_1; UNKNOWN_1.
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SEQUENCE 217
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Yamazaki N., Ueshima R., Terrett J.A.,
Segawa R., Kobayashi T., Numachi K., Ue
Watanabe K., Thomas R.H.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-1021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          J. Mol. Evol. [2]
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"Complete DNA sequence of the mitochondrial genome (Gastropoda: Pulmonata).";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=382;
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NCBI_TaxID-28835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             71 LLLVTLAL 78
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ilarity 100.0%;
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    27353 MW;
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0; Mismatches
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    302E0EA668861EE1 CRC64;
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eda T., Nishikawa
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kawa K.
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Best Local S
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InterPro; IPR001726; Glyco_hydro_19.
Pfam; PP00187; Chitin_bind_1; 1.
Pfam; PP00182; Glyco_hydro_19; 1.
PRINTS; PR00451; CHITINBINDNG.
PR000574; Glyco_hydro_19; 1.
SMART; SM00270; ChtBD1; 1.
PROSITE; PS00773; CHITINASE_19_1; 1.
PROSITE; PS00774; CHITINASE_19_2; 1.
PROSITE; PS000726; CHITINASE_19_2; 1.
PROSITE; PS000726; CHITINASE_19_2; 1.
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Q9HYU9;
01-MAR-2001
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065330;
SEQUENCE FROM N.A.
STRAIN-ATCC 15692 / PAO1;
MEDILINE-20437337; PubMed-10984043;
Stover C.K., Pham X.-Q.T., Erwin A.L., Mizoguchi S.D., Warrener
Stover G.K., Brinkman F.S.L., Hufnagle W.O., Kowalik D.J., Lagro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TISSUE-ROOT NODULE;
Kim H.-B., An C.-S.;
"Isolation and characterization of cDNA clone encoding acidic chitinase from the root nodule of Elaeagnus umbellata.";
Submitted (APR-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF061805; ARC16010.1; -.
HSSP; P23951; 2BAA.
HSSP; P23951; 2BAA.
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Eukaryota; Viridiplantae; Streptophyta; Enbryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosideurosids I; Rosales; Elaeagnaceae; Elaeagnus.
                                                                                                                                                                                                    Pseudomonas aeruginosa.
Bacteria; Proteobacteria;
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01-AUG-1998 (TrEMBLrel. 07, Last sequence up
01-UN-2002 (TrEMBLrel. 21, Last annotation
Accidic chitinase (EC 3.2.1.14)
                                                                                                                                                                                                                                                                         Hypothetical
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                                                                                                                                                                                 Pseudomonas
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O8WMS2;
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            SEQUENCE FROM N.A.
TISSUE-LACRIMAL GLAND;
Lind-Ayres M.R., Crow J.M., St
Lind-Ayres M.R., Crow J.M., St
Remington S.G.;
"Lipophilin AL2 of the male re
Submitted (NOV-2001) to the EN
EMBL; AY063770; AAL40859.1; --
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Zhao C., Nguyen T.X., Lehrer R.I.;
Zhao C., Nguyen T.X., Lehrer R.I.;
"Rabbit Lipophilins.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ EMBL; AF308614; AAG42802.1; ...
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
SMART; SM00096; UTG; 1.
                                                                                                                                                                                                                                                                            Lipophilin AL2.
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Ve.
Mammalia; Eutheria; Lagomorpha; Leporidae;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eu
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus
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01-MAR-2001 (TrEMBLrel.
01-MAR-2002 (TrEMBLrel.
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EMBL, AE004751; AAG06682.1; -.
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01-DEC-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
01-JUN-2002 (TrEMBLrel. 2
"Genome seventh biology."; science 282:2012-2018(1998). EMBL; AL132952; CAC70119.1; EMBL; AL132952; MA: 12323 MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Caenorhabditis elegans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Y51H4A.26 protein.
Y51H4A.26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Globodera pallida.
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                                                                                                                                                                                         SEQUENCE FROM N.A. MEDLINE-99069613;
                                                                                                                                                                                                                                                                                  Submitted
                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=6239;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Rhabditidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Peloderinae;
                                                                                                                                                                                         PubMed=9851916;
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Nematoda;

oda; Chromadorea; Caenorhabditis.

Rhabditida; Rhabditoidea;

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the EMBL/GenBank/DDBJ

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766B3FC05D9C14E0 CRC64;

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Q9BH17;
Q1-JUN-2001 (TrEMBLrel. 17,
Q1-JUN-2001 (TrEMBLrel. 17,
Q1-DEC-2001 (TrEMBLrel. 19,
EMRFamilde-related peptide 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            InterPro; IPR000329; Uteroglobin. Pfam; PF01099; Uteroglobin; 1. SEQUENCE 96 AA; 10510 MW; 48E
                                                                                                                                                                                                                                                                   MEDLINE-21414099; PubMed-11522352;
Kimber M.J., Fleming C.C., Bjourson A.J., Halton D.W.,
"FMRFamide-related peptides in potato cyst nematodes.";
MOI. Blochem, Parasitol, 116:199-208(2001).

EMBL; AJ300488; CAC32452.1; -.

EMBL; AJ300488; CAC32452.1; -.

EMBL; BJ 98 FMRFAMIDE-RELATED PEPTIDE
CCHAIN 89 98 FMRFAMIDE-RELATED PEPTIDE
                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Nematoda;
Tylenchoidea; Heteroderidae;
NCBI_TaxID=36090;
                                                                                                                                                                                   Similarity
7; Conserv
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                                                       PRELIMINARY;
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21,
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Last annotation updat
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24;
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; Globodera.
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Best Local
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Best Local
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01-MAY-1997 (TrEMBLrel. 03, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Hypothetical 14.4 kDa protein.
T16A1.5.
                                                                                                                                                                                                                                                                                                                                                                                                                                   P91445
P91445;
01-MAY-1997
01-MAY-1997
01-DEC-2001
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Q91XL7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a Hibernating Mammal.";
Submitted (JUN-2001) to the EMBL/GenBar
EMBL; AF395869; ARK7258.1; .

InterPro; IPR001981; Colipase.
Pfam; PF02740; Colipase; 1.

Pfam; PF02740; Colipase; 1.

PROSITE; PS00121; COLIPASE; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE-PANCREAS;
Squire T.L., Bauer V.W., Lowe
"Genomic Organization of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-2001 (TremBLrel. 19, 01-DEC-2001 (TremBLrel. 19, 01-MAR-2002 (TremBLrel. 20, Pancreatic colipase.
                                                                                                                                   "Genome sequence of the nematode C. elinvestigating biology. The C. elegans Science 282:2012-2018(1998).
                                                                                                                                                                                                                     SEQUENCE FROM N.A. STRAIN=BRISTOL N2; MEDLINE=99069613;
                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                      Caenorhabditis elegans.
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NCBI_TaxID=43179;
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
          "The sequence of C. elegans Submitted (JAN-1997) to the
                                                                                                                                                                                                                                                                                                        NCBI_TaxID=6239
                                                                                                                                                                                                                                                                                                                                Rhabditidae;
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                                                    Du 2.,
                                                                        STRAIN-BRISTOL N2;
                                                                                              SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                              Peloderinae;
                                                                                                                                                                                                                       PubMed=9851916;
                                                                                                                                                                                                                                                                                                                            Nematoda; Chromadorea; Rhabditida; Rhabditoidea; cinae; Caenorhabditis.
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cosmid T16A1.";
EMBL/GenBank/DDBJ databases
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Last sequence up
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Pred. No.
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Pancreatic Triacylglycerol Lipase
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Craniata; Vertebrata; Euteleostomi;
Sciurognathi; Sciuridae; Sciurinae;
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; Mismatches
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ns Sequencing
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25;
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5. 26;
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Best Local
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Q985Y1;
Q1-OCT-2001 (
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Q1-MAR-2002 (
Hypothetical I
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Q9EX03;
01-MAR-2001
01-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN-A3(2) / M145;
Bentley S.D., Chater K.F., Cerdeno-Tarraga A.-M., Challis G.L.,
Thomson N.R., James K.D., Harris D.E., Quail M.A., Kieser H.,
Harper D., Bateman A., Brown S., Chandra 3., Chen C.W., Collins M.,
Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,
Huang C.-H., Kieser T., Larke L., Murphy L., Oliver K., O'Neil S.,
Rabbinowitsch E., Rajandream M.A., Rutherford K., Rutter S.,
Seeger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K.
Warren T., Wietzorrek A., Woodward J., Barrell B.G., Parkhill J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Streptomyces coelicolor.

Bacteria; Firmicutes; Actinobacteria; Actinobacteridae;
Actinomycetales; Streptomycineae; Streptomyces.
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01-MAR-2001 (TrEMBLrel. 16, Last sequence up
01-JUN-2002 (TrEMBLrel. 21, Last annotation
Hypothetical protein SCO1123.
SCO1123 OR 2SCG38.16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (JUL-2001) to the EMBL/GenBank/IDBJ databases: EMBL; U80454; AAB37873.1; -. Hypothetical protein. SEQUENCE 123 AA; 14402 MW; D9E9DB1374:C4DF2 CRC64;
                  Bacteria; Proteobacteria; alpha subdivis!on; Rhizobiaceae group;
Phyllobacteriaceae; Mesorhizobium.
                                                                  Rhizobium loti (Mesorhizobium loti)
                                                                                         MLL7476.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A
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SEQUENCE FROM N.A.
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STRAIN-MAFF303099;
MEDLINE-21082930; pubMed-11214968;
MEDLINE-21082930; pubMed-11214968;
Matcanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T.,
Matcanabe A., Idesawa K., Ishikawa A., Kawashima K., Kimura T.,
Kishida Y., Kiyokawa C., Kohara M., Matsumoto M., Matsuno A.,
Mochizuki Y., Nakayama S., Nakazaki N., Shimpo S., Sugimoto M.,
Takeuchi C., Yamada M., Tabata S.;
"Complete genome structure of the nitrogen-fixing symbiotic bacterium Mesorhizobium loti.";
DNA Res. 7:331-338(2000).
EMBL, AP003011; BAB53572.1;
EMBL, AP003011; BAB53572.1;
                                                                                                                                                                                                                                                                                                                                                                       "Genome sequence of the hyperthermophilic crenarchaeon Pyrobaculum aerophilium.";
Proc. Natl. Acad. Sci. U.S.A. 99:984-989(2002).
EMBL; AE009916; AAL64741.1; -.
Heat shock; Complete proteome.
SEQUENCE 139 AA; 14562 MW; 96BE53856A3BE60F CRC64;
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STRAIN=IM2 / ATCC 51768 / DSM 7523;
PubMed=11792869;
Fitz-Gibbon S.T., Ladner H., Kim U.-J., Stetter K.O., Simon M.I.,
Miller J.H.;
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01-MAR-2002 (TrEMBLrel. 20, Created)

01-MAR-2002 (TrEMBLrel. 20, Last sequence update)

01-MAR-2002 (TrEMBLrel. 20, Last annotation update)

Possible heat shock protein, htpX homolog part 1, authentic
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Hypothetical protein; Complete proteome.
SEQUENCE 133 AA; 14765 MW; 7EBFD066159E231A CRC64;
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Archaea; Crenarchaeota; Thermoprotei; Thermoproteales; Thermoproteaceae; Pyrobaculum.
NCBI_TaxID=13773;
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LENGTH: 90 AMINO ACIDS
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NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 32:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: 201-994-1700
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz TITLE OF INVENTION: Human Endometrial Specific Statistic OF INVENTION: Binding Factor I, II and III
                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
TOPOLOGY: LINEAR
                                                                                                                                                   61
                                                                                                                                                                                 61
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STATE: NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN ADDRESSEE: CECCHI, STEWART & OLSTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                             MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                               GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                            MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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                                                                                                                                                                                                                                                                                                                                                                                                        AMINO ACID
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     Cohen, Maurice
                                                                                                                                                                                                                                                                               Conservative
                  Billing-Medel, Patricia A.
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                                                                                                                                                                                                                                                                                         100.0%;
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                                                                                                                                                                                                                                                                             0;
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Pred. No. 3.3e-49;
; Mismatches 0;
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                                                                                                                                                                                                                                                                                                        Length 90;
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                                                                                                                                              US-09-583-169-4
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                                                                                            Sequence 4, Application US/09583169 Patent No. 6338948 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                        Best Loc
Matches
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 847/335-1729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for
CURRENT APPLICATION DATA
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ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                              APPLICANT: Jian Ni, TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 90 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION: NAME: Becker, Cheryl L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA: APPLICATION NUMBER:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Le
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TITLE OF INVENTION: REAGENTS AND MITTIFULE OF INVENTION: FOR DETECTING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                CORRESPONDENCE ADDRESS:
                               NUMBER OF SEQUENCES:
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                                                                                                                                                                                                            61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                            61
                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                        MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                     MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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Russell, John C.
Stroupe, Steven D.
Stroupe, Steven D.
WENTION: REAGENTS AND METHODS USEFUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hodges, Steven C. Klass, Michael R.
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                                              Binding
                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                           Guo-Liang Yu and Reiner Gentz
Human Endometrial Specific St
Binding Factor I, II and III
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BYRNE, BAIN, GILFILLAN,
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                                                                                                                                                                                                                                                                                                                                     Score 450; DB 4;
Pred. No. 3.3e-49;
0; Mismatches 0;
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Steroid-

Length 90;

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Gaps

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US-09-215-818-6
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; ORGANISM: Homo Sapiens
US-09-215-818-6
                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: COlpitts, Tracey
TITLE OF INVENTION: REAGENTS AND METHODS USEFUL FOR
TITLE OF INVENTION: DETECTING DISEASES OF THE BREAST
FILE REFERENCE: 5972.US.P2
                                                  SEQ ID NO 6
LENGTH: 90
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Best Local Similarity 100.0%;
Matches 90; Conservative
                                                                                                                                                                                                                                                                                                                                                                                         Sequence 6, Application US/09215818A Patent No. 6379671
                                                                                                                                                                       EARLIER APPLICATION NUMBER: 08/912,276
EARLIER FILING DATE: 1997-08-17
EARLIER APPLICATION NUMBER: 08/697,105
EARLIER FILING DATE: 1996-08-19
EARLIER FILING DATE: 1996-08-19
EARLIER FILING DATE: 1996-08-19
                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/215,818A CURRENT FILING DATE: 1998-12-18
                                                                                     NUMBER OF SEQ ID NOS: 6 SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                      EARLIER FILING DATE: 1996-08-19
                                                                                                                                       EARLIER FILING DATE: 1997-08-15
EARLIER APPLICATION NUMBER: 08/697,106
                                 LENGTH: 90
TYPE: PRT
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LENGTH: 90 AMINO ACIDS
TYPE: AMINO ACID
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MEDIUM TYPE: 3.5 INCH DISKETTE
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OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
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NEW JERSEY
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Pred. No. 3.3e-49;
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; MOLECULE TYPE: US-08-912-276-23
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GENERAL INFORMATION:
Query Match 76.0%; Score 342; DB 4; Best Local Similarity 100.0%; Pred. Nc. 8e-36; Matches 69; Conservative 0; Mismatches
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                                                                                                                                                          INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: FastSEQ for Windows Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                           SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
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                                                                                                LENGTH: 69 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                             NAME: Becker, Cheryl L
REGISTRATION NUMBER: 35,441
REFERENCE/DOCKET NUMBER: 59
                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM CONTROL OPERATING SYSTEM:
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                                                                                    TOPOLOGY:
                                                                                                                                                                                       TELEFAX: 847/938-2623
                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90;
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60064-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Abbott Park
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08912276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100 Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stroupe, Steven D.
REAGENTS AND METHODS USEFUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hodges, Steven C. Klass, Michael R. Kratochvil, Jon D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Granados,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gordon, Julian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Colpitts,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cohen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Billing-Medel, Patricia A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Friedman,
                                                                                                                                                                                                                                                                                                                                                                                                                 IBM Compatible
                                                                                      linear
                                                                       No. 6183952e
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                                                                                                                                                                                                                                                                                                                                                            US/08/912,276
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Pred. No. 3.3e-49;
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                            DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              THE BREAST
                            Length 69;
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US-09-263-810-2

Sequence 2, Application US/09263810

Patent No. 6174992

GENERAL INFORMATION:
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                                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/014,724
FILING DATE: MARCH 21, 1996
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                          STRANDEDNESS:
TOPOLOGY: LII
MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKE
COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                              REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 32:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific Steroid-
TITLE OF INVENTION: Binding Factor I, II and III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/821,451A FILING DATE: March 21, 1997
                                                                                                       61
                                                                                                                                                                                                                                                                                                                                          TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: ROSELAND
STATE: NEW JERSEY
                                                                                                                                 61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: CARELLA, BYRNE, BAIN, GILFILLAN, ADDRESSEE: CECCHI, STEWART & OLSTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61
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                                                                                                                                                           COUNTRY:
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                                                                                                    VKILKKCSV 69
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                                                                                                                                                                                                                                                                                                 PROTEIN
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                                                                                                                                                                                                                       61.6%; Score 277; DB 3; 58.9%; Pred. No. 1.5e-27; tive 17; Mismatches 20
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                                                                                                                                                                                                                        20;
                                                                                                                                                                                                                                                    Length 90
                                                                                                                                                                                                                       Indels
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Best Local
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US-09-263-810-2
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                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09583169 Patent No. 6338948
                                                                                                                                                                                                                                             GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz
TITLE OF INVENTION: Human Endometrial Specific Sta
TITLE OF INVENTION: Binding Factor I, II and III
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 201-99
INFORMATION FOR SEQ
ZIP: 07068
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 32:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-994-1700
TELEPAX: 201-994-1744
                                                                                            STREET: OF CITY: ROSELAND CITY: NEW JERSEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: L. MOLECULE TYPE:
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LENGTH: 90 AMINO ACIDS
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz TITLE OF INVENTION: Human Endometrial Specific Steroid-TITLE OF INVENTION: Binding Factor I, II and III NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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STATE: NEW JERSEY
                                                                                                                                                                          ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                              EVKHCTDQISFKKRLSLEKVLVEIVKKCGV 90
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58.9%;
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& OLSTEIN
                                                                                                                                                                                                                                                      Specific Steroid-
II and III
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; TOPOLOGY: LINEAR
; MOLECULE TYPE: PROTEIN
US-09-583-169-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 10
US-08-912-276-22
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TELEFAX: 201-994-1744
INFORMATION FOR SEQ ID NO: 2:
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APPLICATION NUMBER: 08/821,451
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION NUMBER: US/08/912,276
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                    APPLICANT: Roberts-Rapp, Lisa
APPLICANT: Russell, John C.
APPLICANT: Stroupe, Seren D.
APPLICANT: STROUGH, SEAGENTS AND METHODS USEFUL
TITLE OF INVENTION: FOR DETECTING DISEASES OF THE BREAST
NUMBER OF SEQUENCES: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                          ADDRESSEE: Abbott Laboratories
STREET: 100 Abbott Park Road
CITY: Abbott Park
STATE: IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match 61.6%; Score 277; DB 4; Length 90;
Local Similarity 58.9%; Pred. No. 1.5e-27;
nes 53; Conservative 17; Mismatches 20; Indels
                                                                                                                                                    COUNTRY: USA
ZIP: 60064-3500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                Granados, Edward N.
Hodges, Steven C.
Klass, Michael R.
Kratochvil, Jon D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cohen, Maurice
Colpitts, Tracey L.
Friedman, Paula N.
Gordon, Julian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Billing-Medel, Patricia A.
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ATTORNEY/AGENT INFORMATION: INAME: Becker, Cheryl L REGISTRATION UNMBER: 35,4 REFERENCE/DOCKET NUMBER: 35,4 REGISTRANDEDNESS: single ropology: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: linear NOLECULE TYPE: No. 6183952e US-08-912-276-22 Query Match POPOLOGY: LINEAR NOLECULE TYPE: POPOLOGY: LINEAR NOLECULE TYPE: POPOTEIN	
E: Becker, Cheryl L. ESTRATION NUMBER: 35,441 ESTRATION NUMBER: 35,441 ESTRATION NUMBER: 35,441 ESTRATION NUMBER: 3972.US.P1 COMMUNICATION INFORMATION: EPHONE: 647,935-1729 ERAX: 847,935-1729 ERAX: 847,938-2623 EX. MCE CHARACTERISTICS: ESTRATION COLUMBER: 30,232 ESTRATION COLUMBER: 30,232 ESTRATION COLUMBER: 30,232 ASINCALARD ANDENESS: 19191 ASINCALARD ANDENESS: 2101 LAKPDAPPEAVAAKLEVKCIDOMSLOKESLIAEV-UKILKKCSV 90 IIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIIII	ON: ION DATA: NUMBER: INFORMATION: r, Cheryl L RUMBER: 35,441

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; MOLECULE TYPE:
US-08-821-451A-26
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US-08-821-451A-26
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                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                  Matches
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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itent No. 6066724
                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/014,724
APPLICATION NUMBER: 60/014,724
FILING DATE: March 21, 1996
ATTORNEY/AGENT INFORMATION:
NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: 3.5 INC
COMPUTER: IBM PS/2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                             LENGTH: 90 AMINO ACIDS
TYPE: AMINO ACID
STRANDEDNESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/821,451A FILING DATE: March 21, 1997
60 LEVKRCVDQMSNGDRLVVAETLVYIFLECGV 90
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                                                             1 MKLSVCLLLVTLALCCYQANA-EFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAK 59
                    LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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NEW JERSEY
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E: CECCHI, STEWART & OLSTEIN
6 BECKER FARM ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jian Ni, Guo-Liang Yu and Reiner Gentz
IVENTION: Human Endometrial Specific St
VENTION: Binding Factor I, II and III
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Conservative
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                                                                                                                              44.9%; score 202; DB 3;
48.4%; Pred. No. 3.6e-18;
ive 15; Mismatches 30
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US-09-263-810-26
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US-09-263-810-25
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                                                               GENERAL INFORMATION:
APPLICANT: Jian Ni, Guo-
TITLE OF INVENTION: Huma
TITLE OF INVENTION: Bind
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                           Patent No. 6174992
                                                                                                                                                                                               Sequence
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SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
TYPE: AMINO ACID
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APPLICATION NUMBER:
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SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5 INCH DISKETTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
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TOPOLOGY: LIN
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              STREET:
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 CITY:
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                                                                                                                                                                                                                                                                                                                                                                                           1 MKLSVCLLLVTLALCCYQANA-EFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAK 59
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30, 6174992
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              6 BECKER FARM ROAD
                                                                                                    Jian Ni, Guo-Liang Yu and Reiner Gentz
WYENTION: Human Endometrial Specific St
WYENTION: Binding Factor I, II and III
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                               CARELLA, BYRNE, BAIN, GILFILLAN, CECCHI, STEWART & OLSTEIN
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Pred. No. 3.6e-18;
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                                                                                                    Specific Steroid-
II and III
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STATE:

NEW JERSEY

ZIP: 07068
COMPUTER READABLE FORM:

COUNTRY: USA

MEDIUM TYPE: 3.5 INCH DISKETTE COMPUTER: IBM PS/2

WORD PERFECT 5.1

MS-DOS

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; MOLECULE TYPE:
US-09-263-810-26
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Patent No. 6338948
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 26:
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SOFTWARE: WORD PERFECT
CURRENT APPLICATION DATA:
                                                                                                                        COMPUTER: IBM PS/2
OPERATING SYSTEM: MS-DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH DISKETTE
COMPUTER: IBM PS/2
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LENGTH: 90 AMINO ACIDS
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NAME: MULLINS, J.G.
REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
               PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jian Ni, Guo-Liang Yu and Reiner Gentz TITLE OF INVENTION: Human Endometrial Specific Steroid-TITLE OF INVENTION: Binding Factor I, II and III NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: 201-994-1700
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APPLICATION NUMBER: (
FILING DATE:
ATTORNEY/AGENT INFORMATION:
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                                                                  APPLICATION NUMBER: US/09/583,169
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                     STREET: 6 BECKER I
CITY: ROSELAND
STATE: NEW JERSEY
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                                08/821,451
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Search completed: January 2, 2003, 14:55:38 Job time: 15 secs
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 90 AMINO ACIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 33,073
REFERENCE/DOCKET NUMBER: 325800-521 (PF257)
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                         TYPE: AMINO ACID
STRANDEDNESS: SIN
                                                                              60 LEVKRCVDQMSNGDRLVVAETLVYIFLECGV 90
                                                                                                     60 LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                                                                                      1 MKLSVCLLLVTLALCCYQANA-EFCPALVSELLDFFI'ISEPLFKLSLAKFDAPPEAVAAK 59
1 IELSLCLLIM-LAVCCYEANASQICELVAHETISFLIKSEEELKKELEMYNAPPAAVEAK 59
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Perfect score:
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Word size :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Scoring table:
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                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   seq length: 0 seq length: 2000000000
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/cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/US10_PUBCOMB.pep:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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          0 US-09-825-301-77

0 US-09-110-716-29

0 US-09-34-054-1

0 US-09-985-911-4

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0 US-09-110-716-37

0 US-09-110-716-38

0 US-09-989-738-260

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0 US-09-989-730-260

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 117078
Sequence 77, Appl Sequence 29, Appl Sequence 1, Appli Sequence 4, Appli Sequence 37, Appl Sequence 37, Appl Sequence 260, App Sequence 260, App
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61	ш ш	Query M Best Lo Matches	SULT 1 Sequence Sequence Sequence APPLIC TITLE FILE CURREN		20 20 22 22 22 23 24 24 25 26 27 27 28 28 28 28 28 28 28 28 28 28 28 28 28
GVKRCTD	MKLSVCI MKLSVCI	Match Local Simi es 90;	-301-77 e 77, App. No. USZON. INFORMAT: ANT: Hougl ANT: Yu, ANT: Yu, ANT: Yu, ANT: Yu, TEPERENCE:		
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GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90	CPALVSELLDFF:	; Score 90; DB 10; Lengt ; Pred. No. 3.8e-81; 0; Mismatch:s 0; Inde	ITIONS AND KITS FOR OF BREAST CANCER 25,301 .	ALIGNMENTS	10 US-09-989-731-260 10 US-09-989-732-260 10 US-09-991-732-260 10 US-09-991-732-260 10 US-09-991-73-260 10 US-09-993-604-260 10 US-09-993-604-260 10 US-09-985-911-260 10 US-09-985-911-260 10 US-09-985-911-260 10 US-09-985-911-260 10 US-09-985-911-242-5162 10 US-09-912-242-5162 10 US-09-925-297-617 10 US-09-925-297-617 10 US-09-912-022-293 10 US-09-912-022-293 10 US-09-912-031-393-43 10 US-09-935-291-998
	ISEPLFKLSLAKFDAPPEAVAAKL 60	th 90; lels 0; Gaps	THE DETECTION		Sequence 260, App Sequence 27, Appli Sequence 27, Appli Sequence 210, App Sequence 210, App Sequence 27, Appli Sequence 28, Appli Sequence 29, Appli Sequence 29, Appli Sequence 210, Appli Sequence 213, Appli Sequence 213, Appli Sequence 214, Appli Sequence 214, Appli Sequence 214, Appli Sequence 214, Appli Sequence 215, Appli Sequence 214, Appli Sequence 2168, Appli Sequence 39434, A Sequence 36472, A Sequence 36472, A
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RESULT 2
US-09-110-716-29
; Sequence 29, Application US/09110716A
; Patent No. US20020034739A1
; GENERAL INFORMATION:

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61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90

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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 29
LENGTH: 90
TYPE: PRT
ORGANISM: lipophilin B
US-09-110-716-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Lehrer, Robert I.
APPLICANT: Zhao, Chengquan
APPLICANT: Clasgow, Benjamin J.
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF CERTAIN TUMORS
FILE REFERENCE: 22000-20596.00
CURRENT FILLING DATE: 1998-07-07
CURRENT FILLING DATE: 1998-07-07
CURRENT FILLING DATE: 1998-07-07
COMMERC OF SEQ ID NOS: 41
                                                                                                                                     TELEPHONE: (650) 855-05
TELEFAX: (650) 845-4166
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                    COMDUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/934,054
FILING DATE: 21-Aug-2001
PRIOR APPLICATION DATA:
                                                                                                                                                                                           APPLICATION NUMBER: US/08/747,547
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0077 US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 94025-6936
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MURRY, Lynn E.
Goll, Surya K.
Goll, Surya K.
Hawkins, Phillip R.
TITLE OF INVENTION: BREAST TUMOR SPECIFIC PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Akerblom, Ingrid E.
Hillman, Jennifer L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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US20020107385A1
                                                      TYPE: amino acid
STRANDEDNESS: double
                                   TOPOLOGY: linear
                                                                                             LENGTH: 90 amino acids
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STATE: CA
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TYPE: <Unknown>
DESCRIPTION: SE
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SEQ ID
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NO:
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; TYPE: PRT
; ORGANISM: LpnB
US-09-110-716-37
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US-09-110-716-37
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PRIOR APPLICATION NUMBER: 09/263,810
PRIOR FILING DATE: 1999-03-08
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR APPLICATION NUMBER: 08/821,451
PRIOR FILING DATE: 1997-03-21
PRIOR FILING DATE: 1996-03-21
PRIOR FILING DATE: 1996-03-21
                                                                    SOFTWARE: PatentIn SEQ ID NO 37
                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                              Sequence 37, Application US/09110716A Patent No. US20020034739A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US Patent No. US20020151012A1
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                                                                                                                                                   APPLICANT: Lehrer, Robert I.
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF
FILE REFERENCE: 22000-20596.00
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best
                                                                                                   CURRENT APPLICATION NUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
NUMBER OF SEQ ID NOS: 41
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TITLE OF INVENTION: HUMAN ENDOMETRIAL SPECIFIC STEROID-BINDING FACTOR I, II AND FILE REFERENCE: PF257D3
CURRENT APPLICATION NUMBER: US/09/985,911
CURRENT FILING DATE: 2001-11-06
CURRENT FILING DATE: 2001-11-06
PRIOR APPLICATION NUMBER: 09/83,169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 90
TYPE: PRT
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nes 90; Conserv
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ilarity 100.0%;
Conservative (
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Pred. No. 3.8e-81;
D; Mismatches 0;
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Pred. No. 3.8e-81;
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                                                                                                   SOFTWARE: Annomax
SEQ ID NO 44240
LENGTH: 50
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Best Local
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                                                                                                                                                               PRIOR APPLICATION NUMBER: US 09/774,203
PRIOR FILING DATE: 2001-01-29
NUMBER OF SEQ ID NOS: 49117
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
PRIOR FILING DATE: 2000-06-30
                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: PCT/US01/00670 PRIOR FILING DATE: 2001-01-30
                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2001-0: PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2001-0:
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PRIOR APPLICATION NUMBER: PCT/US01/00662
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PRIOR APPLICATION NUMBER: PCT/US01/00663
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PRIOR APPLICATION NUMBER: PCT/
PRIOR APPLICATION 2001-01-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Penn, Sharron G. APPLICANT: Rank, David R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/US01/00665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE:
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                                      LENGTH: 50
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: MAP TO AC004127.1 OTHER INFORMATION: EXPRESSED IN LUNG,
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Hanzel, David K.
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Pred. No.
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                                                                                                                                           Engine vers. 1.1
  SIGNAL =
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9.9e-61;
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US-09-992-598-260
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US-09-110-716-38
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LENGTH: 74
TYPE: PRT
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Best Local s
Matches 8
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Matches
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APPLICANT: Zhao, Chengquan
APPLICANT: Zhao, Chengquan
APPLICANT: Glasgow, Benjamin J.
TITLE OF INVENTION: PEPTIDES CHARACTERISTIC OF CERTAIN TUMORS
FILE REFERENCE: 22000-20596.00
CURRENT FILING DATE: 1998-07-07
RUMBER: US/09/110,716A
CURRENT FILING DATE: 1998-07-07
RUMBER OF SEQ ID NOS: 41
                                                   APPLICANT:
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APPLICANT:
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OTHER INFORMATION:
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8; Conserv
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                                                      Gurney, Austin L.
Kljavin, Ivar J.
                                                                                                         Godowski,Paul J.
Grimaldi,J.Christopher
                                                                                                                                                                         Goddard, Audrey
                                                                                                                                                                                                 Gerritsen, Mary E.
                                                                                                                                                                                                                                  Gerber, Hanspeter
                                                                                                                                                                                                                                                             Fong, Sherman
                                                                                                                                                                                                                                                                                        Ferrara, Napoleone
                                                                                                                                                                                                                                                                                                                      Eaton, Dan L.
                                                                                                                                                                                                                                                                                                                                               Desnoyers, Luc
                                                                                                                                                                                                                                                                                                                                                                             Botstein, David
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Pan, James
                        Napier, Mary A.
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larity 100.0%;
Conservative
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EXPRESSED IN BRAIN, SIGHAL = 0.95
EXPRESSED IN PLACENTA, SIGNAL = 1.2
EXPRESSED IN BONE MARROH, SIGNAL = 1.1
EXPRESSED IN ADULT LIVE!, SIGNAL = 1.4
EXPRESSED IN ADULT LIVE!, SIGNAL = 1.2
EXPRESSED IN BONE MARROH = 1
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100.0%; Pr
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APPLICANT APPLICANT: APPLICANT

Tumas,

Daniel

Colin K.

Roy, Margaret Ann c+ewart, Timothy A.

Paoni, Nicholas F.

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PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR APPLICATION NUMBER: 60/06250
PRIOR APPLICATION NUMBER: 60/065186
PRIOR APPLICATION NUMBER: 60/065311
PRIOR APPLICATION NUMBER: 60/065311
PRIOR APPLICATION NUMBER: 60/06770
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
PRIOR APPLICATION NUMBER: 60/078910
PRIOR APPLICATION NUMBER: 60/08332
PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/08322
PRIOR APPLICATION NUMBER: 60/08726
PRIOR APPLICATION NUMBER: 60/087106
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APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C20
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OR APPLICATION NUMBER: 60/088033

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OR APPLICATION NUMBER: 60/088212

OR FILING DATE: 1998-06-05

OR APPLICATION NUMBER: 60/088217

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PR APPLICATION NUMBER: 60/089947
DR FILING DATE: 1998-06-19
DR APPLICATION NUMBER: 60/089948
DR FILING DATE: 1998-06-19
DR APPLICATION NUMBER: 60/089952
DR FILING DATE: 1998-06-19
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DR APPLICATION NUMBER: 60/090246
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OR FILING DATE: 1998-06-10
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NUMBER: 60/089598: 1998-06-17 60/089599

NUMBER: 60/089653 60/089907 60/089801 60/089600

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NUMBER: 60/090252

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                 APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C66
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OR APPLICATION NUMBER: 60/091360
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APPLICATION NUMBER: 60/090696
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Publication No. US20
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          PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
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                                                                                   CURRENT APPLICATION NUMBER: US/09/989,735
CURRENT FILING DATE: 2001-11-19
                                                                                                                  APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C61
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APPLICATION NUMBER: 60/092182
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APPLICATION NUMBER: 60/091626
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APPLICATION WINDER: 60/091633
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Grimaldi, J.Christopher
Gurney, Austin L.
Kijavin, Ivar J.
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Tumas, Da
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Gerritsen, Mary E.
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PRIOR FILING DATE: 1998-06-16 PRIOR APPLICATION NUMBER: 60/089532 PRIOR APPLICATION NUMBER: 60/089538 PRIOR APPLICATION NUMBER: 60/089598 PRIOR FILING DATE: 1998-06-17 PRIOR APPLICATION NUMBER: 60/089599 PRIOR FILING DATE: 1998-06-17 PRIOR APPLICATION NUMBER: 60/089599 PRIOR FILING DATE: 1998-06-17 PRIOR APPLICATION NUMBER: 60/089600 PRIOR FILING DATE: 1998-06-17 PRIOR APPLICATION NUMBER: 60/089601 PRIOR APPLICATION NUMBER: 60/089908 PRIOR FILING DATE: 1998-06-18 PRIOR APPLICATION NUMBER: 60/089909 PRIOR FILING DATE: 1998-06-19 PRIOR APPLICATION NUMBER: 60/089909 PRIOR FILING DATE: 1998-06-19 PRIOR APPLICATION NUMBER: 60/089909 PRIOR APPLICATION NUMBER: 60/089947 PRIOR APPLICATION NUMBER: 60/089947 PRIOR APPLICATION NUMBER: 60/089947 PRIOR APPLICATION NUMBER: 60/090246 PRIOR FILING DATE: 1998-06-29 PRIOR APPLICATION NUMBER: 60/090252 PRIOR FILING DATE: 1998-06-22 PRIOR APPLICATION NUMBER: 60/090252 PRIOR APPLICATION NUMBER: 60/090254 PRIOR APPLICATION NUMBER: 60/090254 PRIOR APPLICATION NUMBER: 60/090443 PRIOR APPLICATION NUMBER: 60/090443 PRIOR APPLICATION NUMBER: 60/090444 PRIOR FILING DATE: 1998-06-24 PRIOR APPLICATION NUMBER: 60/090445 PRIOR APPLICATION NUMBER: 60/090557 PRIOR APPLICATION NUMBER: 60/090540 PRIOR APPLICATION NUMBER: 60/090540 PRIOR APPLICATION NUMBER: 60/090540 PRIOR APPLICATION NUMBER: 60/090557 PRIOR APPLICATION NUMBER: 60/090557 PRIOR PRILING DATE: 1998-06-24 PRIOR APPLICATION NUMBER: 60/090567 PRIOR APPLICATION NUMBER: 60/090567 PRIOR APPLICATION NUMBER: 60/090569 PRIOR APPLICATION NUMBER: 60/090569 PRIOR APPLICATION NUMBER: 60/090669 PRIOR APPLICATION NUMBER: 60/			
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OR APPLICATION NUMBER: 60/065311
OR FILING DATE: 1997-11-13
OR APPLICATION NUMBER: 60/066770
OR FILING DATE: 1998-02-25
OR APPLICATION NUMBER: 60/078910
OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/08332
OR APPLICATION NUMBER: 60/08322
OR FILING DATE: 1998-03-20
OR APPLICATION NUMBER: 60/08760
OR FILING DATE: 1998-05-28
OR APPLICATION NUMBER: 60/088026
OR FILING DATE: 1998-05-02
OR APPLICATION NUMBER: 60/088026
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P2730PlC19
CURRENT APPLICATION NUMBER: US/09/990,444
CURRENT APPLICATION NUMBER: US/09/990,444
CURRENT FILING DATE: 2001-11-14
PRIOR APPLICATION NUMBER: 60/049787
PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/065186
PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
PRIOR APPLICATION NUMBER: 60/06570
PRIOR APPLICATION NUMBER: 60/06570
PRIOR APPLICATION NUMBER: 60/075945
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PRIOR FILING DATE: 1998-02-25
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US-09-990-444-260
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OR FILING DATE: 1998-07-02

OR APPLICATION NUMBER: 60/091633

OR FILING DATE: 1998-07-02

OR APPLICATION NUMBER: 60/091978

OR FILING DATE: 1998-07-07

OR APPLICATION NUMBER: 60/091982

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OR APPLICATION NUMBER: 60/092182

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REFILING DATE: 1998-07-02
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Wood, William I.
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art, Timothy A.
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OR APPLICATION NUMBER: 60/089907
OR FILING DATE: 1998-06-18
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TITLE OF INVENTION: Acids Encoding the same
FILE REFERENCE: P2730P1C69
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OR FILING DATE: 1998-07-02
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APPLICANT: Watanabe, Colin K.
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
APPLICANT: Wood, William I.
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C53
CURRENT APPLICATION NUMBER: US/09/991,181
CURRENT FILING DATE: 1201-11-16
PRIOR APPLICATION NUMBER: 60/049787
PRIOR APPLICATION NUMBER: 60/062250
PRIOR APPLICATION NUMBER: 60/062250
PRIOR APPLICATION NUMBER: 60/065186
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-12
PRIOR APPLICATION NUMBER: 60/065311
PRIOR FILING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/065770
PRIOR FILING DATE: 1997-11-24
PRIOR APPLICATION NUMBER: 60/075945
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DR APPLICATION NUMBER: 60/087609
DR FILING DATE: 1998-06-02
DR APPLICATION NUMBER: 60/087759
DR FILING DATE: 1998-06-02
DR APPLICATION NUMBER: 60/087827
DR APPLICATION NUMBER: 60/087827
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FILING DATE: 1998-02-25
APPLICATION NUMBER: 60/078910
FILING DATE: 1998-03-20
APPLICATION NUMBER: 60/083322
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APPLICATION NUMBER: 60/0
FILING DATE: 1998-05-28
APPLICATION NUMBER: 60/0
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R APPLICATION NUMBER: 60/09
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RR APPLICATION NUMBER: 60/09
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APPLICATION NUMBER: 60/0
FILING DATE: 1998-07-01
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APPLICATION NUMBER: 60/092182
FILING DATE: 1998-07-09
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APPLICATION NUMBER: 60/
FILING DATE: 1998-07-02
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RESULT 15 US-09-993-687-260

PRIOR

FILING DATE:

1998-06-19

APPLICATION NUMBER: 60/ FILING DATE: 1998-06-18 APPLICATION NUMBER: 60/

60/089908

R APPLICATION NUMBER: 60/08
R FILING DATE: 1998-06-19
RR APPLICATION NUMBER: 60/08
RR FILING DATE: 1998-06-19
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60/090246 60/089952 60/089948 60/089947 PRIOR PRIOR PRIOR PRIOR

FILING DATE: 1998-06-18 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-18

60/089907

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60/089801

APPLICATION NUMBER: 60/089599
FILING DATE: 1998-06-17
APPLICATION NUMBER: 60/089600
FILING DATE: 1998-06-17
APPLICATION NUMBER: 60/089653

APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-16 APPLICATION NUMBER: 60/0 FILING DATE: 1998-06-17

60/089514 60/089512

60/089532

APPLICATION NUMBER: 60/1 FILING DATE: 1998-06-16

APPLICATION NUMBER: 60/ FILING DATE: 1998-06-12 APPLICATION NUMBER: 60/ FILING DATE: 1998-06-16

FILING DATE: 1998-06-APPLICATION NUMBER: 60/ FILING DATE: 1998-06-1: APPLICATION NUMBER: 60/

60/089105

60/089440

APPLICATION NUMBER: 60/089538 FILING DATE: 1998-06-17

APPLICATION NUMBER: 60/089598 FILING DATE: 1998-06-17

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GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
APPLICANT: Ashker, Kevin P.
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APPLICANT:
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APPLICANT:
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CURRENT APPLICATION NUMBER: US/09/993,687
CURRENT FILING DATE: 2002-11-14
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APPLICANT: Zhang, Zemin
TITLE OF INVENTION: Secreted and Transmembrane
TITLE OF INVENTION: Acids Encoding the Same
FILE REFERENCE: P2730P1C11
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PRIOR FILING DATE: 1997-06-16
PRIOR APPLICATION NUMBER: 60/062250
PRIOR FILING DATE: 1997-10-17
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OR APPLICATION NUMBER: 60/083322
OR FILING DATE: 1998-04-28
OR APPLICATION NUMBER: 60/084600
OR FILING DATE: 1998-05-07
OR APPLICATION NUMBER: 60/087106
OR FILING DATE: 1998-05-28
OR APPLICATION NUMBER: 60/087607
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/087759
OR FILING DATE: 1998-06-02
OR APPLICATION NUMBER: 60/087759
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OR APPLICATION NUMBER: 60/087759
OR APPLICATION NUMBER: 60/087827
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R APPLICATION NUMBER: 60/06
R APPLICATION NUMBER: 60/07
R FILING DATE: 1997-11-24
R APPLICATION NUMBER: 60/07
R FILING DATE: 1998-02-25
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APPLICATION NUMBER: 60/088029
FILING DATE: 1998-06-04
                                                                                       FILING DATE: 1998-06
APPLICATION NUMBER:
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APPLICATION NUMBER: 60/
FILING DATE: 1998-06-04
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                      APPLICATION NUMBER: 60/088030
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Gurney, Austin L.
Kljavin, Ivar J.
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Godowski, Paul J.
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Botstein, David
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R APPLICATION NUMBER: 60/01

R FILING DATE: 1998-06-22

R APPLICATION NUMBER: 60/05

R FILING DATE: 1998-06-22

R FILING DATE: 1998-06-23

R APPLICATION NUMBER: 60/09

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OR APPLICATION NUMBER: 60/089600

OR FILING DATE: 1998-06-17

OR APPLICATION NUMBER: 60/089653

OR FILING DATE: 1998-06-17

OR APPLICATION NUMBER: 60/089801
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R APPLICATION NUMBER: 6
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APPLICATION 1998-06-16
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APPLICATION NUMBER: 60/089952
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APPLICATION NUMBER: 60/090246
FILING DATE: 1998-06-22
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FILING DATE: 1998-06-10
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FILING DATE: 1998-06-05
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Search completed: January Job time : 12 secs
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PRIOR FILLING DATE: 1998-06-26
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PRIOR APPLICATION NUMBER: 60/091544
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Best Local Similarity 100
Matches 8; Conservative
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PRIOR FILING DATE: 1998-07-07
PRIOR APPLICATION NUMBER: 60/091982
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PRIOR FILING DATE: 1998-06-24
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PRIOR FILING DATE: 1998-06-24
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                                                                                                                                                                                                                                                                                                           Score
                                                                                                                                                                                                                                                                                                                                                                                                                                          is derived by analysis of the total score distribution.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:
2: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:
3: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1982.DAT:
4: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1983.DAT:
4: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1983.DAT:
5: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1984.DAT:
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7: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1987.DAT:
8: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1989.DAT:
8: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1989.DAT:
8: /SIDSZ/gcgdata/geneseq/geneseqp-embl/AA1989.DAT:
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Copyright (c) 1993 - 2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /SIDS2/gcgdata/geneseq/geneseqp-embl/AA2000.DAT:*
/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2001.DAT:*
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21
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AAW35803
AAW59776
AAW59771
AAW89613
AAB13787
AAB07501
AAB07501
AAB07501
AAB07501
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(without alignments)
342.644 Million cell updates/sec
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Compugen Ltd.
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BU101 antigenic pe
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BU101 antigenic pe
Lipophilin B polyp
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	19221 19221
ALIGNMENT:	AAB31681 ABB09634 ABB11907 AAY48606 AAC07521 AAE07522 AAE07524 AAE07524 AAE07527 AAB07527 AAB07527 AAB07527 AAB07527 AAB07527 AAB07509 AAB07676 AAB31680 AAW33628 AAW366704 AAW112391 AAW66704
	An endometrial spe Human breast tumou Human breast tumou Human breast tumou Human lipophilin B Human endometrial spe Amino acid sequenc Human endometrial spe Human endometrial spe Human secreted pro Human 5' EST relat Human screted pro Human fire manmary tis Bovine mammary tis Peptide #7513 enco Human bone marrow Peptide #7665 enco Human profein of human tis Peptide enco Human profeil prote Human profeil prote Human PROB12 prote Membrane-bound pro Human PROB12 polyp Human PROB12 polyp Human PROB12 (NOS BUIO) antigenic pe

Endometrial specific steroid-binding factor II; ESF II; human; Clara cell secretory protein; endometrium; phospholipase A2 inhibitor; polychlorinated biphenyl; antiaggregant; inflammation; asthma; rhinitis; cystic "ibrosis; airway disease; neoplasia; atopy; therapy; diagnosis. 21-MAR-1996; 21-MAR-1996; 25-SEP-1997 Protein Peptide Human endometrial specific steroid-binding factor 27-MAR-1998 AAW35803; AAW35803 standard; Protein; 90 AA. (HUMA-) HUMAN GENOME SCI INC WO9734997-A1 Homo sapiens (first entry) 96WO-US03857. 96WO-US03857 /label= Sig_peptide 22..90 /label= Mat_protein Location/Qualifiers

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RESULT 2
AAW59776
ID XXX ACC XXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence comprises human endometrial specific steroid binding C factor II (ESF II), a protein that inhibits phospholipase A2 activity, binds to polychlorinated biphenyl compounds, reduces C foreign protein antigenicity inhibits monocyte and neutrophil c chemotaxis and phagocytosis, inhibits monocyte and neutrophil c chemotaxis and phagocytosis, inhibits platelet aggregation, regulates elocasnoid levels in the human uterus and controls the growth of endometrial cells. The amino acid sequence was deduced from a cDNA clone (see AAT94831) derived from cyclohaximide-treated CC my cells. ESF I (see AAM35802) and ESF III (see AAM35804) are also claimed. Human ESF II has about 49% identity with rat prostatic steroid-binding protein. Recombinant ESF I, and III can be expressed in host cells for use in claimed methods (a) for treating a patient in need of ESF I, II or III (including expression of the polypeptide in vivo) and (b) for identifying compounds which bind to and inhibit activation of the ESF polypeptide. hESF I, II and III may be used to treat inflammation, asthma, rhinitis, cystic fibrosis, airway disease, neoplasia and atopy.
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                       Akerblom IE,
                                                                             (INCY-) INCYTE PHARM INC
                                                                                                                                        12-NOV-1996;
                                                                                                                                                                                                 07-NOV-1997;
                                                                                                                                                                                                                                                      22-MAY-1998.
                                                                                                                                                                                                                                                                                                                 WO9821331-A1
                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                            antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human steroid-binding protein C1; hSBP1; hSBP2; breast cancer; prot
gene therapy vector; ribozyme; probe; hybridisation; amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Amino acid sequence of the human steroid binding protein Cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW59776;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW59776 standard;
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                       Goli SK,
                                                                                                                                        96US-0747547
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                       Hawkins PR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   90
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Pred. No. 9.2e-51;
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                       Hillman
                       JL,
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                       Murry LE;
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CC This is the amino acid sequence of the human steroid-binding protein CC C1 (hSBP1) used in the method of the invention for the diagnosis, CC monitoring and treatment of breast cancer. HSBP1 and hSBP2 are useful CC as markers for breast cancer, i.e. measuring levels of hSBP1 and hSBP2 cased for diagnosis or monitoring the disease, to identify subjects CC at risk and to discriminate between different forms of cancer for CC selection of appropriate therapies. They may also be used for drug cscreening. Nucleic acids encoding hSBP1 and hSBP2 can be used in gene CC therapy vectors to overexpress the steroid-binding proteins, preventing CC binding of steroids, or antisense sequences, ribozymes. Their nucleic CC acids can also be used for the diagnosis and monitoring (by quantifying CC expression of hSBP), as source of probes for hybridisation and CC amplification of genomic or related sequences for studying regulation of gene function and for mapping the genomic sequence. Antibodies are used CC as diagnostic reagents in standard immunoassays for hSBP.
                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Fig 1; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New human steroid binding diagnosis, monitoring and
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1998-297935/26.
N-PSDB; AAV41579.
90
ĀΑ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proteins C1 and C2 - useful treating breast cancer, and
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RESULT 3
AAW54271
ID AAW5
                                                                                                                                 Matches
                                                                                                                                           Query Match
Best Local
 AAW54271 standard;
                                                  61
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                                                                                                   1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                          ᅟ
                                                 GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                         MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                 90;
                                                                                                                                            Similarity
                                                                                                                                Conservative
 Protein;
                                                                                                                                         100.0%;
  90
                                                                                                                                 0;
                                                                                                                                         Score 450; DB 19;
Pred. No. 9.2e-51;
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26-FEB-1998

WO9807857-A1

Homo sapiens.

BU101; breast immunisation;

cancer; diagnosis;
drug screening; epi

screening; epitope

prevention; treatment; gene therapy;

BU101 antigenic peptide epitope

29-JUL-1998 (first entry

WPI; 1998-169161/15.

Roberts-rapp L,

Billing-medel PA,

Granadosen,

Hodges

Friedman PN; MR, Kratochv

Kratochvil JD;

Russell JC, Cohen M,

Stroupe SD; Colpitts TL, es SC, Klass (ABBO) ABBOTT LAB.

15-AUG-1997; 19-AUG-1996; 19-AUG-1997;

97US-0912276 96US-0697105 97WO-US14665

N-PSDB; AAV26461

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC This represents a BUIO1 polypeptide sequence. BUIO1 is a member of the CC uteroglobin family of proteins and is over-expressed in breast cancer. CC Cells transformed with a recombinant expression system comprising a cC sequence derived from the BUIO1 open reading frame and with at least 50 percent identity to the sequences shown in AAV26458 to AAV26451 are used CC percent identity to the sequences shown in AAV26458 to produce BUIO1 polypeptides containing at least 1 peltope. These are used to detect BUIO1 specific antibodies which are used correspondingly CC to detect BUIO1 antigens. The BUIO1 polynucleotide sequences can be used in a method for detecting the presence of a target BUIO1 polynucleotide. The various assays are used for diagnosis, prognosis, staging, connitoring, treating and preventing diseases of the breast (especially CC cancer and its metastases), and also for determining susceptibility. The BUIO1 polypeptides are also useful in drug screening, e.g. to identify CC antagonists of BUIO1, potentially useful therapeutically and as targets CC antegretically to neutralise BUIO1 polypeptides. Fragments of the BUIO1 nucleic acid are useful as probes and primers, e.g. for detection CC of altered gene expression or in fluorescent in situ hybridisation, also CC genetic immunisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Treatment of endometrial cancer, mammary cancer, endometriosis or endometrial fibroids - comprises administering endometrial steroid
                                                                WPI; 1999-080843/07.
                                                                                                                                                                                                  09-JUN-1998;
                                                                                                                                                                                                                                                                                                                             Endometrial steroid binding protein II; ESBPII; cancer; detection; endometriosis; endometrial fibroid; mammary cancer.
                                                                                                                                                                                                                                                                                                                                                                             Endometrial steroid binding protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW89613 standard; Protein;
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                                                   N-PSDB; AAX00069
                                                                                                 Schmidt CJ,
                                                                                                                                 (SMIK ) SMITHKLINE BEECHAM CORP
                                                                                                                                                                  09-JUN-1997;
                                                                                                                                                                                                                                   17-DEC-1998
                                                                                                                                                                                                                                                                WO9856248-A1
                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                 Wang X;
                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                  97US-0049015.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 450; DB 19;
Pred. No. 9.2e-51;
; Mismatches 0;
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RESULT 5
AAB13787
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Best Local
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                                                                      Multimeric polypeptide antigen and antibody specific useful for diagnosing, detecting and treating breast
                                                                                                                                                                                                                                      Colpitts TL,
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Pred. No. 9.2e-51;
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Claim 1; Page 124; 124pp; English.

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RESULT 6
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        The present sequence represents a BU101 polypeptide. The BU101 gene is transcribed from breast tissue. The specification describes a method detecting the presence of a target BU101 polynucleotide in a test sample. The method comprises contacting the sample with at least one BU101-specific polynucleotide (AAASB875-80), and detecting bound polynucleotides. The method and BU1010 polynucleotides are useful for detecting the presence of BU101 polynucleotides. The methods may be used for the diagnosis of breast disease, indicated by the formation
                                                                                                                                                                 Detecting presence of target BU101 polynucleotide in sample useful detection of breast cancer, comprises contacting sample with BU101-specific polynucleotide and determining binding -
                                                                                                                                         Claim
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No. 9.2e-51;
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cc antiallergic, and cytostatic properties. The polynucleotides are used in cand/or prevent inflammation, asthma, rhinitis, cystic fibrosis, air way classase, neoplasia and atopy. The polynucleotides are also used to complete protein antigenicity, inhibit monocyte and neutrophil chemotaxis and phagocytosis, inhibit platelet aggregation, regulate elocasancid clavels in the human uterus and control the growth of endometrial cells. The polynucleotides are also useful for detecting complementary completed are used to detect complementary polynucleotides are used to attend of a mutated form of hESF I, II and III cassociated with a dysfunction will provide a diagnostic tool that can define diagnosis of a disease or susceptibility to a disease which cresults from under-expression, over-expression or altered expression of hESF I, II and III cassociated with a dysfunction will provide a diagnostic tool that can define diagnostic or a disease which the complementary polynucleotides are used to detect complementary polynucleotides are used to detect complementary polynucleotides such as a second or adjacase which the complementary polynucleotides are used to detect complementary polynucleotides and that can define diagnostic reagent. Detection will provide a diagnostic tool that can define diagnostic or adjacase which the complementary polynucleotides are used to detect complementary polynucleotid
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                                                                                                                                                                                                                                                                                                                                     This invention relates to nucleic acid molecules encoding portions of the human endometrial specific steroid-binding factors I, II, and III. Also included in the invention are hSSF I, II, and III polypeptide sequences. The nucleotide sequence exhibit antiasthmatic, antiinflammatory,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Endometrial specific steroid-binding asthma; rhinitis; cystic fibrosis; a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human endometrial specific steroid-binding factor is useful for treating asthma, rhinitis, cystic
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Pred. No. 9.2e-51;
; Mismatches 0;
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ir way disease; neoplasia; atopy;
is inhibitor; endometrial cancer; s
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Best Local
                                                                                                          The present sequence represents a human endometrial specific steroid-binding protein (ESBP) II. The ESBPII protein is overexpressed in breast tumours. The specification describes a method for diagnosing prostate or a gynaecological cancer. The method comprises measuring levels of ESBPII in cells, tissues or body fluids of a patient, and comparing this to levels from a normal control, where a variance in levels indicates cancer. The method is used to diagnose, stage, monitor, image or treat prostate or gynaecological cancer. The gynaecological cancers include breast, endometrial, ovarian and uterine cancer.
                                                                                                                                                                                                                                                                               Diagnosing, staging, monitoring, imaging and treating prostate and gynaecological cancers by measuring levels of endometrial specific steroid-binding protein (ESBP)II expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; endometrial specific steroid-binding protein II; ESBPII breast tumour; prostate cancer; gynaecological cancer; cancer; endometrial cancer; ovarian cancer; uterine cancer.
                                                                                                                                                                                                                                                      Claim 6; Page 31-32;
                                                                                                                                                                                                                                                                                                                                        WPI; 2000-303648/26.
N-PSDB; AAA14953.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A human endometrial specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                endometrial cancer. They are also useful for chromosome identification. The present sequence represents a hESF \, II protein sequence identified in the invention.
                                                                                   Sequence
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Pred. No. 9.2e-51;
                          Score 450; DB 21;
Pred. No. 9.2e-51;
; Mismatches 0;
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20-JUL-2000;
27-JUL-2000;
18-DEC-2000;
                      from tissue of interest (TI), performing DNA microarray analysis to identify first subset of polynucleoticles (SFI) at least 2-fold over expressed in TI, and performing quantitative polymerase chain reaction (PCR) analysis on SPI to identify second subset of (P). The method is useful for determining the presence on absence of a cancer cell in a patient, monitoring the progression on cancer in a patient using a biological sample such as blood, serum, lymph nodes, bone marrow, sputum, urine or a tumour biopsy sample. The methods are useful for determining the presence or absence of or monitoring progression of prostate, breast, colon, ovarian, lung, head and neck, lymphoma, leukemia, melanoma, liver, gastric, kidney, bladder, pancreatic or endometrial cancer. The present sequence represents the lipophilin B polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genetic cancer;
                                                                                                                                                                                                                                                                                       Identifying tissue (tumour)-specific pylynucleotides overexpressed in tissue of interest as compared to control tissue, for detecting cancer cells in patient, comprises DNA microarray analysis or quantitative
                                                                                                                                                                                                 The invention relates to identifying tissue-specific polynucleotides (P) that involves performing a genetic sultraction to identify pool of (P)
                                                                                                                                                                                                                                                                           cells in patient, comprises polymerase chain reaction -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lipophilin B
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                                                                                                                                                                                                                                                                                                                                                                                                                       (CORI-) CORIXA CORP.
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 90
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                                                                                                                                                                                                                                                                                    Query Match
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Matches 90; Conser
                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a complex comprising a lipophilin-like polypeptide linked by at least one disulphide bond to a second lipophilin-like polypeptide. Lipophilin-like protein are members of uteroglobin superfamily. Lipophilin-like proteins are useful in the preparation of vaccines. The complex containing lipophilin-like proteins are useful for treating or preventing breast, ovarian or prostate cancer. The complex is also used for determining the presence or absence of cancer in a patient, or monitor the progress of cancer in a patient. Lipophilin DNA is also useful in gene therapy. The present sequence is human lipophilin B protein.
 AAB31681;
                                     AAB31681 standard;
                                                                                                                                                                                                                                                                                                                                                             Sequence
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28-JUN-2000; 2000US-0215735
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; Page 72; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 prostate cancer
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                                                                                                                                                                                                                                                                                Score 450; DB 22;
Pred. No. 9.2e-51;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                       90;
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RESULT 12
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a human endometrial specific steroid binding factor (hESF). The specification describes hESF1, hESF11, and hESF111. hESF11. It and III polypeptides, and polypucleotides encoding them are useful for treating and preventing inflammation, asthma, rhinitis, cystic fibrosis, airway disease, neoplasia and atopy, inhibiting phospholipase A2 activity, binding polychlorinated biphenyls, reducing foreign protein antigenicity, inhibiting monocyte and neutrophil chemotaxis and phagocytosis, inhibiting platelet aggregation, regulating eicosanoid levels in the human uterus, and for controlling the growth of endometrial cells. hESF polypeptides and nucleotides are also useful for research, biological, clinical or the property of the province of the provin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New human endometrial specific steroid binding factors, treating and preventing inflammation, asthma, rhinitis, fibrosis, airway disease, neoplasia and atopy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1; Fig 2; 36pp; English.
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21-MAR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      An endometrial specific steroid binding factor
ABB09634 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapeutic purposes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-JAN-2001.
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                                                                                                                                                                                                                                                  . 61
                                                                                                                                                                                                                                      GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
                                                                                                                                                                                                                                                                                                                                             MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                                                                                                                                                                                                                                                                                                                                     MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                                                                                                                                              GVKRCTDQMSLQKRSLIAEVLVKILKKCSV
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DB; AAF25213.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90 AA;
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97US-0821451.
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1..21
Protein;
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90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 450; DB 22;
Pred. No. 9.2e-51;
Mismatches 0;
A
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29-MAY-2002 ABB09634;

(first entry)

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                                                                                                                                                                                                                                                                                          Query Match
Best Local
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              Human; cytokine;
                                     Human breast tumour-associated protein homologue,
                                                               11-JAN-2002
                                                                                                              ABB11907 standard; peptide; 117
                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antibody specific for human endometrial specific steroid-binding factor (hESF) III, useful for detecting hESF III protein in biological sample and to isolate or identify clones expressing the protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ni J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-MAR-1996;
21-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human endometrial specific steroid-binding factor (hESF)
 haematopoiesis regulation;
                                                                                                                                                                                                                                                                                                                                                 cnromatography. Agonists and antagonists of hESF proteins are useful for treating and/or preventing susceptibility to asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN
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                                                                                                                                                                                                                                                    بر
                                                                                                                                                                          GVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
                                                                                                                                                                                                                                                   MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2002-215019/27.
                                                                                                                                                                                                                            MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFTSEPLFKLSLAKFDAPPEAVAAKL 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Yu G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        endometrial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABL41782.
                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       dometrial specific steroid-binding factor;
steroid-binding protein; hESF I; hESF II;
                                                                                                                                                                                                                                                                                                                             90
                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fig 1;
                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-0583169
                                                                                                                                                                                                                                                                                                                             AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-014724P.
97US-0821451.
99US-0263810.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "signal peptide" 22..90
cell proliferation; cell
egulation; tissue growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ₽,
                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "mature protein"
                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                        Score 450; DB 23; Pred. No. 9.2e-51;
                                                                                                                                                                                                                                                                             Mismatches
              differentiation; growth factor;
 immunomodulator;
                                     SEQ ID
                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                   Length
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 activin;
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haematopolesis regulatory activity; tissue growth activity;

chamunomodulatory activity; activity or inhibin-related activities;

chemotactic or chemokinetic activities; haemostatic, thrombotic or

chemotactic or chemokinetic activities; haemostatic, thrombotic or

cthrombolytic activities; receptor or i.gand activities; or may be

cinvolved in oncogenesis, cancer cell proliferation or metastasis.

Ce pepending on their biological activities, polypeptides and nucleotides of

the invention are useful for preventing, treating or ameliorating medical

conditions, e.g., by protein or gene therapy. Such conditions include

cancers, haematopoietic disorders (e.g., myeloid or lymphoid cell

conditions, e.g., by protein or gene therapy. Such conditions include

cancers, haematopoietic disorders (e.g., myeloid or lymphoid cell

conditions, e.g., by protein or gene therapy. Such conditions in arthritis),

proliferative retinopathy, atherosclerssis, coronary heart disease,

conditions are useful or promote wound

conditions activities involved with tissue regeneration and

conditions activities may be used to promote wound

conditions activities may be used in the treatment of viral,

bacterial and fungal infections in addition to immune disorders.
                                         Polypeptides with growth factor activity may be used in cell cultures to promote cell growth. For example, such polypeptides may be used to manipulate stem cells in culture to give rise no neuroepithelial cells that can be used to augment or replace cells damaged by illness, autoimmune disease or accidental damage. The polypeptides and nucleotides may also be used in the diagnosis of the above conditions, and in drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             or polypeptides in a sample, and methods of identifying compounds which bind to polypeptides of the invention. Although novel, many of the polypeptides of the invention have homology to known proteins, thereby giving an insight into their probable biological activities, and hence potential therapeutic applications. The polypeptides of the invention me have various activities, including cytckine, cell proliferation or cell differentiation activities; stem cell growth factor activity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequences ABB10981-ABB12330 represent 1:50 novel human polypeptides, and sequences ABA0825-ABA09574 represent nucleic acids encoding them. The invention also relates to vectors and recombinant host cells comprising a nucleotide of the invention, methods of producing the novel polypeptides, antibodies against the polypeptides, methods of detecting the nucleotides of antibodies against the polypeptides, methods of detecting the nucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     myeloid cell disorder; lymphoid cell disorder; asthma; arthritis; chronic inflammatory condition; proliferative retinopathy; atherosclerosis; coronary heart disease; arterial ischaemia; bone disorder; osteoporosis; vascular growth disorder; tissue regeneration; wound healing; infection; immune disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 20; Page 275; 1963pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human proteins and DNA encoding sequences useful for preventing, treating or ameliorating a medical condition in a mammalian subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
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27-APR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HYSE-) HYSEQ INC
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proliferation; metastasis; cancer; tumour haematopoietic disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001-457740/49
DB; ABA09151.
                         techniques.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               osteopathic; vasotropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-0496914.
2000US-0560875.
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                       The present sequence
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                       represents a
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                                                                                                                                This invention decribes novel human nucleic acid sequences from tumor breast tissue which have cytostatic activity. The nucleic acid sequences can be used to produce and isolate full-length gene sequences. They can be used to express proteins, which can be used as tools to find an activity against breast cancer. The sequences can be used in sense or antisense form. They are especially useful for medicaments for gene therapy to treat breast cancer. AAY48540-Y48617 represent protein
                                                                                                                                                                                                                      Claim 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                       Sequence
                                                                                                                                                                                                                                       Human nucleic acid sequences and tissue, useful for breast cancer
                                                                                                                                                                                                                                                                                                Specht T,
                                                                                                                                                                                                                                                                                                                                                                                            DE19813839-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                Expressed sequence tag; EST; human; breast; cancer; gene therapy;
treatment; tumour; cytostatic; medicament.
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                                                                                                                          the invention.
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                          MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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DB; AAZ33666.
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Pred. No. 1.3e-50;
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Pred. No. 1.3e-50;
Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                               presence or absence of cancer in a patient, or monitor the progress of cancer in a patient. Lipophilin DNA is also useful in gene thera. The present sequence is human lipophilin B S11 3 3 protein.
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28-JUN-2000;
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Result
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Perfect score:
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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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    hits satisfying chosen parameters:
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Copyright (c) 1993 - 2003 Compugen
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CYA7_BOVIN
FLAE_METUA
ME10_CAEEL
PSC3_RAT
VATA_CHLPN
D11_DICDI
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ALB1_SALSA
ALB2_SALSA
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UGR1_HUMAN
CYTF_MOUSE
IRA2_YEAST
SY04_CHICK
FEL2_FELCA
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lepus capen
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                                                                                                                                                                                                                                                                                                                                                                                                                        homo sapien
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MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES. SUBCELLULAR LOCATION: Secreted (Potential). TISSUE SPECIFICITY: HIGHEST EXPRESSION MAS FOUND IN SKELETAL MUSCLE. EXPRESSED AS WELL IN THYMUS, TRACHEA, KIDNEY, STEROII RESPONSIVE TISSUES (PROSTATE, TESTIE, UTERUS, BREAST AND OVAL AND SALIVARY GLAND.

BREAST AND OVARY),

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ESTRAMUSTINE,	FUNCTION: MAY BIND	2 2	gu)	167	FROM N.A.		NCBI_TaxID=9606;	Mammalia; Eutheria;	Eukaryota; Metazoa;	Homo sapiens (Human).	SCGB1D2 OR LIPHB.	Lipophilin B precursor					_				_	_	۰	_	_	_	_	_	_	_	_	_		
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A CHEMOTHERAPEUTIC	ANDROGENS AND OTHER	"Lipophilins: human peptides homologous to rat prostatein.";	Zhao C., Nguyen T., Yusifov T., Glasgow	MEDLINE-99167354; PubMed-10066439;				Primates;	Chordata;			(Secretoglobin family	Last	Last	Created)						1	-	_	_	_	_	ب	_	_	1	_	_		
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PROSTATE CANCER	MAY ALSO BIND							٠	Euteleostomi;												pse	gallus	archaeoglob	escherichia	ric	hon	homo	homo	poc.	rat	met	eps		
CE	₹								٠.												pseudomonas	lus	hae	hei	rickettsia	homo sapien	ō	ō	podospora	rattus norv	methanococc	epstein-bar		•
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문 Qy Query Match Best Local S Matches 90 SMART; SM00096; UTG; 1 Signal. 1 23 CHAIL 22 90 EMBL; AJ224172; CAA11864.1; -. Genew; HGNC:18396; SCGB1D2. Genew; HGNC:18396; Uteroglobin. InterPro; IPR000329; Uteroglobin; 1. Pfam; PF01099; Uteroglobin; 1. PRINTS; PR00486; UTEROGLOBIN. This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (see http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch). SEQUENCE SIMILARITY: BELONGS TO THE UTEROGLOFIN FAMILY. LIPOPHILIN SUBFAMILY. Similarity 1 22 90 AA; Conservative 21 90 9925 100.0%; MW. 0; POTENTIAL. LIPOPHILIN B. 17BB555ED035D1AF CRC64; Score 450; DB 1; Pred. No. 6.7e-45; ; Mismatches 0; Length Indels 0; 0

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LPPA_HUMAN
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InterPro; IPRO00329; Uteroglobin; 1.
Pfam; PF01039; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
SMART; SM00096; UTG; 1.
                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                           SEQUENCE
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15-JUN-2002 (Rel. 41, Last annotation update)
Lipophilin A precursor (Secretoglobin family 1D member SCGB1D1 OR LIPHA.
                                                                                                                                                                                                                                                                                   EMBL; AJ224171; CAA11863.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Glasgow B.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE OF 22-90, MEDLINE-98385871;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. MEDLINE-99167354; PubMed-10066439;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'Lipophilin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ehrer R.I., Xu G.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                 POPHILIA a novel heterodimeric protein of human tears.";
IS Lett, 432:163-167(1998).
IS Lett, 432:163-167(1998).
FUNCTION: MAY BIND ANDROGENS AND OTHER STEROIDS, MAY ALSO BIND ESTRANUSTINE, A CHEMOTHERAPEUTIC AGENT USED FOR PROSTATE CANCER. MAY BE UNDER TRANSCRIPTIONAL REGULATION OF STEROID HORMONES.
SUBUNIT: HETERODIMER OF A LIPOPHILIN A AND A LIPOPHILIN C (MAMMAGLOBIN B) MONOMER ASSOCIATED HEAD TO HEAD.
TISSUE SPECIFICITY: EXPRESSED IN LACHRYMAL GLAND, THYMUS, KIDNEY TESTIS, OVARY AND SALLYARY GLAND.

MASS SPECTROMETRY: MM-7574.69; METHOD-Electrospray; RANGE-22-90.
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
                                                                                                                                                                                                                                                                                                                                                                                                                                SUBFAMILY
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                                                        MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
                                          MRLSVCLLLLTLALCCYRANAVVCQALGSEITGFLLAGKPVFKFQLAKFKAPLEAVAAKM
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9898 MW;
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                                                                                                               59.3%;
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Pred. No. 6.8e-24;
                                                                                                                                                        LIPOPHILIN A.
DF2D4F7565A87D34 CRC64;
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MEDINE-80075873; PubMed-7198120;
Liao S., Chen C., Huang I.-Y.;
"Prostate alpha-protein. Complete amino acid
"Prostate alpha-protein nuclear retention of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parker M.C
                                                                                                                                                                                                                                                                                                                                                                                                                                          Delaey B., Rombauts W., Volckeert G., Peeters B., "Identification of a complementary-DNA clone conto sequence information for the C-1-polypeptide of reprotein.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peeters B., Heyns W., Mous J., Rombauts W.; "Structural studies on rat prostatic binding structure of component Cl from subunit F."; Eur. J. Biochem. 123:55-62(1982).
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21-JUL-1986 (Rel. 01, Created)
13-AUG-1987 (Rel. 05, Last sequenc
16-OCT-2001 (Rel. 40, Last annotat
Prostatic steroid-binding protein
                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "The nucleotide sequence of cDNA complementary rat prostatic binding protein."; Eur. J. Biochem. 133:645-649(1983).
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Rombauts W.;
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[2]
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MEDLLNE-82220075; PubMed-6896362;
Parker M.G., Needham M., White R.;
"Prostatic steroid binding protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                            CHAIL SOC. Trans. 10:51-51(1982).

FUNCTION: PART OF PROSTATEIN WHICH IS THE MAJOR SECRETORY
GLYCOPROTEIN OF VENTRAL PROSTATE GLAND.

SUBUNIT: PROSTATEIN IS COMPOSED OF THREE DIFFERENT PEPTIDES
C1, C2 AND C3. THESE FORM COVALENT C1:C3 (F) AND C2:C3 (S)
HETERODIMERS WHOSE NONCOVALENT ASSOCIATION FORMS TETRAMERIC
(C1:C3/C3:C2) PROSTATEIN MOLECULES.

SUBCELLULAR LOCATION: SECRETED.

MISCELLANEOUS: THE HETERODIMER CAN BIND NON-POLAR STEROIDS,
CHOLESTEROL AND A GROUP OF SMALL PROLINE-RICH PEPTIDES.
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                                                                                                                 SUBFAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            м.G.;
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40, Last annotation update)
-binding protein Cl chain precursor
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Rodentia;
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Sciurognathi; Muridae;
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                                                                                                               MEDLINE-83209619; PubMed-6343081; Peeters B., Heyns W., Mous J., Rombauts W.; Peeters B., Heyns W., Mous J., Rombauts W.; "Structural studies on rat prostatic binding structure of component C2 from subunit S."; Eur. J. Biochem. 132:669-679(1983).
                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
MEDLINE-87146484; PubMed-2881277;
Delaey B., Dirckx L., Decourt J.-
Rombauts W.;
                                                                                                                                                                                                                                                                                                                                                 Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10116;
[1]
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01-JAN-1988 (Rel. 06, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Prostatic_steroid-binding protein C2 chain precursor (Prostatein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR000329; Uteroglobin
Pfam; PF01099; Uteroglobin; 1.
SMART; SM00096; UTG; 1.
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EMBL; J00774; AAA41969.1;
EMBL; J00773; AAA41969.1;
EMBL; V01545; CAA24787.1;
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PIR; S42615; S42615.
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                  binding
                                                   Parker
                                                                   MEDLINE=82220075;
                                                                                SEQUENCE OF 1-100 FROM N.A
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                                  "Prostatic steroid
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K -> E (IN REF. 1; AAA41969).
A -> S (IN REF. 6).
CC -> GG (IN REF. 6).
CC -> GG (IN REF. 5).
N -> D (IN REF. 5).
                              protein:
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Pred. No. 8.4e-17;
                                                                                                                                                                                                                                                                                                                                                                              Sciurognathi; Muridae;
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Best Local
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01-OCT-1989 (Rel. 12, Created)
01-OCT-1989 (Rel. 12, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Clara cell phospholipid-binding protein precu-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CONFLICT
SEQUENCE
                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
                 SEQUENCE FROM
                                                NCBI_TaxID=9606;
                                                                                                                 SCGB1A1 OR UGB
                                                                                                                                                                                                                                               UTER_HUMAN
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(Human) о_к

Primates; Chordata;

Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.

CC10

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CCSP (CC10)

protein)

Orotein precursor (CCPBP) (Clara (Utero lobin) (Urine protein 1)

cells

STANDARD;

PRT;

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EMBL; X05034; CAA28708.1; EMBL; V01256; CAA24599.1; EMBL; J00776; AAA51641.1; PIR; A03251; BORT2. PIR; A03671; A26671.
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DISULFID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               use by non-profit institutions as lone as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                           Steroid-binding;
                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR000329; Uteroglobin.
PROSITE; PS00403; UTEROGLOBIN_1;
PROSITE: PS00404; UTEROGLOBIN_2;
                                      1 MKLSVCLLLVTLALCCYQAN-----AEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEA
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FYM: LINKED BY THREE DISULFIDE BONDS TO C3.

MISCELLANEOUS: THE HETERODIMER CAN BIND NON-POLAR STEROIDS.

CHOLESTEROL AND A GROUP OF SMALL PROLINE-RICH PEPTIDES.

SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. LIPOPHILIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GLYCOPROTEIN OF VENTRAL PROSTATE GLAN).
SUBUNIT: PROSTATEIN IS COMPOSED OF THEE DIFFERENT PEPTIDES
C1, C2 AND C3. THESE FORM COVALENT C1:C3 (F) AND C2:C3 (S)
HETERODIMERS WHOSE NONCOVALENT ASSOCIATION FORMS TETRAMERIC
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(C1:C3/C3:C2) PROSTATEIN MOLECULES
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INTERCHAIN (WITH C3) (PROBABLE).
INTERCHAIN (WITH C3) (PROBABLE).
INTERCHAIN (WITH C3) (PROBABLE).
MISSING (IN REF. 3).
I -> T (IN REF. 3).
I -> T (IN REF. 3).
                                                                               13;
                                                                                               Score 171; DB 1
Pred. No. 8.8e-1
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DA65A6A82E677864 CRC64;
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-, identified proteins in human nasal lawage fluid non-smokers and smokers using two-dimensional gel electropeptide mass fingerprinting ";

Proteomics 2:112-120(2002).

(*RAY CRYST***
                                       This
                                                                                                                 "Structure of a human Clara cell phospholipid-binding complex at 1.9-A resolution.";
Nat. Struct. Biol. 1:538-545(1994).
-i- FUNCTION: BINDS PHOSPHATIDYLCHOLINE, PHOSPHATIDYLI POLYCHLORINATED BIPHENYLS (PCB) AND WEAKLY PROGEST INHIBITOR OF PHOSPHOLIPASE A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-93016476; PubMed-1400743; Okutani R., Itoh Y., Hirata H., Kasahara T., Mukaida N., Kawai T.; "Simple and high-yield purification of urine protein 1 using immunoaffinity chromatography; evidence for the identity of urine protein 1 and human Clara cell 10-kilodalton protein.";
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MEDLINE-93250776; PubMed-1284526;
Wolf M. Klug J. Hackenberg R., (
Beato M., Suske G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein."
Biochim. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Human CC10, the homologue of rabbit uteroglobin: chromosomal localization and expression in endomet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Human J.G., Danel C., Chu C., Crystal R.G.;
"Human CC10 gene expression in airway epithelium
locus suggest linkage to airway disease.";
Am. J. Physiol. 268:L565-L575(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Singh G., Katyal S.L., Brown W.I Anthony J., Squeglia N.; "Amino-acid and cDNA nucleotide
                                                                                                                                                                                                                           X-RAY CRYSTALLOGRAPHY (1.
MEDLINE-95393197; PubMed-
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MEDLINE-95250987; PubMed-7733299;
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                                                                                                                                                                                                  Pletcher J., Sax M.;
                                                                                                                                                                                                                  Umland
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chromatogr. A 577:25-35(1992).
                                                            SUBUNIT: Homodimer; antiparallel disulfide-linked TISSUE SPECIFICITY: CLARA CELLS (NONCILIATED CELL: SURFACE EPITHELIUM OF THE PULMONARY AIRWAYS). SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
        SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation - European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                     Ther J., Sax M.:
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                                                                                                                                                                                                                                                                                           electrophoresis
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trial cell lines.";
the EMBL OLL.
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RESULT 6
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Best Local.S
Matches 26
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                             21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
Uteroglobin precursor (Blastokinin).
SCGBIAI OR UGB OR UGL.
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                                                                                                                                SEQUENCE FROM N.A.
MEDLINE-#83290960; PubMed-6309802;
Bailly A., Atger M., Atger P., Ce
Logeat F., Milgrom E.;
"The rabbit uteroglobin gene. Str
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; X13197; CAA31584.1; -.
EMBL; U01101; AAA81885.1; -.
EMBL; U01102; AAA18297.1; -.
EMBL; BC004481; AAH04481.1; -.
EMBL; X59875; CAA42532.1; -.
SEQUENCE FROM N.A. MEDLINE=83220783; Suske G., Wenz M.,
                                                                                       progesterone receptor.";
J. Biol. Chem. 258:10384-10389(1983)
                                                                                                                                                                                                                                                                                   Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF01099; Uteroglobin; 1. PRINTS; PR00486; UTEROGLOBIN. ProDom; PD012475; Uteroglon_sub;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and foentities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                 NCBI_TaxID=9986;
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InterPro; IPR000329; Uteroglobin.
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S26651; S26651.
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  PubMed=6304644;
, Cato A.C.B., E
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INTERCHAIN (WITH C-90').
INTERCHAIN (WITH C-24').
INTERCHAIN (WITH C-24').
R -> G (IN DBSNP:1802634).
/FTId=VAR_012045.
T -> A (IN DBSNP:1802632).
/FTId=VAR_012046.
C -> E (IN REF. 7).
C -> E (IN REF. 7).
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Pred. No. 7.1e
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    Beato
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[11]
SEQUENCE OF 39-77 FROM N.A.
SEQUENCE OF 39-77 FROM N.A.
MEDLINE-81021016; PubMed-7417250;
Chandra T., Woo S.L.C., Bullock D.W.;
Cloning of the rabbit uteroglobin structural gene.";
"Cloning of the rabbit of the structural gene.";
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(8)
SEQUENCE OF 22-91.
SEQUENCE OF 22-91.
MEDLINE-79074850; PubMed-281700;
Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
Popp R.A., Foresman K.R., Wise L.D., Daniel J.C. Jr.;
Pamino acid sequence of a progesterone-binding protein.";
"amino acid sequence of u.S.A. 75:5516-5519(1978).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Suske G., Menne C., Cato A., Wenz M., Beato M. "Characterization and sequence analysis of int DNA sequences transcribed in X.laevis embryos. Prog. Clin. Biol. Res. 85:139-146(1982).
                                                                                                                                 SEQUENCE OF 53-72 FROM N....

MEDLINE-80241888; PubMed-6156676;

MEDLINE-80241888; PubMed-6156676;

Atter M., Perricaudet M., Tiollais P., Milgrom E.;

Bacterial cloning of the rabbit uteroglobin structural

"Anniva Res. Commun. 93:1082-1088(1980).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Popp R.A., Foresman K.R., Wise I Submitted (OCT-1982) to the PIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ponsting1 H., Nieto A., Beato M.; "Amino acid sequence of progesterone-induced Biochemistry 17:3908-3912(1978).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-79187160; PubMed-571719;
Atger M., Mercier J.-C., Haze G., Fri
"N-terminal sequences of uteroglobin
Blochem. J. 177:985-988(1979).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIINE-83157105; PubMed-6299663;
Chandra T., Bullock D.W., Woo S.L.C.;
"Hormonally regulated mammalian gene expression: steady-state level
and nucleotide sequence of rabbit uteroglobin mRNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Isolation and structure of the gene protein uteroglobin.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "The uteroglobin gene region: hormonal regulation, repetitive elements and complete nucleotide sequence of the gene."; Nucleic Acids Res. 11:2257-2271(1983).
                                                                                  MEDLINE=89199637;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      nucleotide
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MEDLINE=79042086; PubMed=568483;
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                     1.64-A resolution.
                                                               Bally R., Delettre J.;
                                                                                                            K-RAY CRYSTALLOGRAPHY
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                                            "Structure and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1:19-26(1981).
  Biol.
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                                                                                                                                                                                                                                                                                                                                                                                                                            tructure of rabbit lung sequence of a cDNA."; 193:247-249(1985).
206:153-170(1989)
                                        refinement of the
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"rnemann J., Wenz M.,
"agne for
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ProDom; PD012475; Uteroglbn_sub; 1.
SMART; SM00096; UTG; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Morize I., Surcouf E., Vaney M.C., Epelboin Y., Buehner M., Fridlansky F., Milgrom E., Mornon J.-P.; "Refinement of the C222(1) crystal form cf oxidized uterogl 1.34-A resolution.";
                               SEQUENCE
                                          HELIX
                                                                                                                           CONFLICT
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                                                                                                                                                           CONFLICT
                                                                                                                                                                                                  CHAIN
                                                                                                                                                                                                             SIGNAL
                                                                                                                                                                                                                       Phospholipase
                                                                                                                                                                                                                                PROSITE; PS00403; PROSITE; PS00404;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=94297152; PubMed=8025221;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRUCTURE BY NMR OF 39-68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Conformation and molecular dynamics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        [mprota S., Pastore A., Mammi
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                                                                                                                                                                                                                                                                                                                                              X K01657; AAA31497.1;

J J00688; AAA31495.1;

J J00688; AAA31495.1;

X X01423; CAA25669.1;

J M32012; AAA31500.1;

J M25090; AAA31500.1;

J M27564; AAA31496.1;

J M25057; AAA31498.1;

M25038; AAA31499.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SECRETED BY THE UTERUS SIMILARITY: BELONGS TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: Homodimer; antiparallel dis INDUCTION: UTEROGLOBIN, SYNTHESIZED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PHOSPHOLIPASE A2
                                                                                                                                                                                                                                                                                                                 A03249; UGRB.
1UTG; 15-OCT-92.
2UTG; 15-OCT-89.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the EN
                                                                                                                                                                                                                                                                                                                                                                                                                                                     s requires a license agreement (see http://www.isb-sib.ch/announce/an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biol. 194:725-739(1987).
                                                                                                                                                                                                                             PS00403; UTEROGLOBIN_1;
PS00404; UTEROGLOBIN_2;
                                                   Ã2
                                                                                                                                                                                                                       inhibitor;
                               9983 MW;
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22.7%;
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                                                                                                                         INTERCHAIR (WITH C-90').
INTERCHAIR (WITH C-24').
T -> F (IN REF. 6).
C -> G (IN REF. 6).
L -> V (IN REF. 5).
DS -> NT (IN REF. 12).
E -> Q (IN REF. 7).
                                                                                                                                                                                                                      Steroid-binding; Signal; 3D-structure.
Score 10:; DB 1; Pred. No. 5.9e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        s.
                               OC1978AAI:5D550CA CRC64;
                                                                                                                                                                                                  UTEROGLOB (N
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lcs calculations
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ZED IN THE UTERUS
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                                                                                                                                                            STRAIN-129/Svy TISSUE-rung;
STRAIN-129/Svy TISSUE-rung;
MEDLINE-94071937; PubMed-7916613;
Ray M.K., Magdaleno S., O'Malley B.W., Demayo F.J.;
"Cloning and characterization of the mouse Clara cell specific 10
"Cloning and characterization of the 5'-flanking region with the human
                                                                                                                                                                                                                                                                 Singh G., Katyal S.L., Brown W.E., Kennedy A.L.; "Mouse Clara cell 10-Kba (CC10) protein: cDNA nucleotide sequence molecular basis for the variation in progesterone binding of CC10 from different species."; cxp. Lung Res. 19:67-75(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Cloning and tissue-specific expression of the cDNA for the mouse clara cell 10 kD protein: comparison of endogenous expression to rabbit uteroglobin promoter-driven transgene expression."; Am. J. Respir. Cell Mol. Biol. 9:231-238(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
Clara cell phospholipid-binding protein precursor (CCPBP) (Clara cells
10 kDa secretory protein) (CCl0) (Uteroglobin) (PCB-binding protein)
(Clara cell 17 kDa protein).
SCGBIAI OR UGB OR CC10 OR UTG.
                                                                                                                                          Biochem. Biophys. Res. Commun. 197:163-171(1993) -1- FUNCTION: BINDS PHOSPHATIDYLCHOLINE, PHOSPHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
STRAIN-129/SV; TIS
                                                                                                                                                                                                                                                                                                                    MEDLINE=93178380; PubMed=8440203; Singh G., Katyal S.L., Brown W.E.
                                                                                                                                                                                                                                                                                                                                                                                             gene."
                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE-94292183; PubMed-8020953; Stripp B.R., Huffman J.A., Bohinski R.J.;
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Margraf L.R., Finegold M.J., Stanley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q06318;
01-JUN-1994
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                                                                                                                                                                                                                                                                                                                                                                                                            "Structure and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Demayo F.J.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=10090;
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                          INDUCTION: BY GLUCOCORTICOIDS.
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                      DEVELOPMENTAL STAGE: APPEARS ON IN THE AIRWAY EPITHELIUM. INDUCTION: BY GLUCOCORTICOIDS.
                                                             SUBUNIT: Homodimer; antiparallel disulfide-linked. TISSUE SPECIFICITY: CLARA CELLS (NONCILIATED CELLS SURFACE EPITHELIUM OF THE PULMONARY AIRMAYS).

DEVELOPMENTAL STAGE: APPEARS ON THE EIGHTEENTH DAY
                                                                                                               POLYCHLORINATED BIPHENYLS (PCB) AND WEAKLY PROGESTERONE, INHIBITOR OF PHOSPHOLIPASE A2.
   SWISS-PROT
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Sciurognathi;
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UTER_LEPCA
ID UTER_L
AC P06913
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DT 01-FEB
DT 15-JUN
DE UTEROG
GN SCGB1A
OS Lepus
OC Mammal
OX NCB1_T
RN [1]
RP SEQUEN
RC TISSUE
RX MEDLIN
RC TOPEZ
RT "Nucle
RY PIC-UT
RL Bioche
CC -!- SI
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Best Local S
Matches 24
                                                                                                                                                     TISSUE-Lung;
MEDLINE-86323069; PubMed-3019311;
Lopez de Haro M.S., Nieto A.;
"Nucleotide and derived amino acid
pre-uteroglobin from the lung of th
Blochem. J. 235:895-898(1986).
-1- FUNCTION: UTEROGLOBIN BINDS PRO
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EMBL; L24372; AAA65446.1; -.
EMBL; X67702; CAAA7936.1; -.
EMBL; U01247; AAA16141.1; -.
PIR; S24783; S24783.
PIR; A53025; A53025.
HSSP; P17559; IUTR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               UTER_LEPCA
P06913;
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     -1- SUBUNIT: Homodimer; antiparallel disulfide-linked.
-1- INDUCTION: UTEROGLOBIN, SYNTHESIZED IN THE UTERUS AND SECRETED BY THE UTERUS UPON INDUCTION BY PROGESTERONE.
-1- SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                           01-JAN-1988 (Rel. 06, Created)
01-FEB-1996 (Rel. 33, Last sequence up
15-JUN-2002 (Rel. 41, Last annotation
Uteroglobin precursor (Blastokinin).
SCGB1A1 OR UGB OR UGL.
Lepus capensis (Brown hare).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
ProDom; PD012475; Uteroglobn_sub;
SMART; SM00096; UTG; 1.
                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrai
Mammalia; Eutheria; Lagomorpha; Leporidae; Lepus
NCBI_TaxID-9981;
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SIGNAL 1 19 BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00403; UTEROGLOBIN_1; 1. PROSITE; PS00404; UTEROGLOBIN_2; 1.
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the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by ar
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InterPro; IPR003628; Uteroglbn_sub.
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                                                                                                              FUNCTION: UTEROGLOBIN BINDS PROGESTERONE SPECIFICALLY AND HIGH AFFINITY. IT MAY REGULATE PROGESTERONE CONCENTRATIONS REACHING THE BLATOCYST. IT IS ALSO A POTENT INHIBITOR OF
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Pred. No. 8
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INTERCHAIN (WITH C-90') (BY SIMILARITY).
INTERCHAIN (WITH C-24') (BY SIMILARITY).
21103A8839497E6C CRC64;
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RESULT 9
UTER_RAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DISULFID
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   UTER_RAT STANDARD; PRT; 96 AA.
p17559;
01-AUG-1990 (Rel. 15, Created)
01-NOV-1990 (Rel. 16, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
16-JUN-2002 (Rel. 41, Last annotation update)
10-NOV-1990 (Clara cells
10-NOV-1990 (Clara cells
10-NOV-1990 (Rel. 41, Last sequence update)
10-NOV-1990 (Rel. 41, Last sequence update)
10-NOV-1990 (Rel. 41, Last sequence update)
10-NOV-1990 (Rel. 15, Created)
10-NOV-1990 (Rel. 15, Created)
10-NOV-1990 (Rel. 16, Last sequence update)
10-NOV
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  SEQUENCE OF 1-18 F
MEDLINE=90272398;
Hagen G., Wolf M.,
                                                                                                                                                                       MEDLINE=90324266; PubMed=2115524;
Nordlund-Moeller L., Andersson O.,
Gillner M., Gustafsson J.-A., Lund
                                                                                                                                                                                                                                                               nucleotide sequences and developmental expression. Prog. Respir. Res. 25:29-35(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      InterPro; IPR003628; Uteroglbh_sub.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
                                                                                                          polychlorinated biphenyls. progesterone-binding protei
                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. Katyal S.L., Singh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRINTS; PR00486; UTEROGLOBIN. ProDom; PD012475; Uteroglbn_s
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; M25609; AAA30960.1; -
PIR; A23825; UGRBL.
                                                                                                                                                     "Cloning,
                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                          Wong-Chong M.-L.;
"Clara cell secretory (10 kDaltons)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DISULFID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00403; UTEROGLOBIN_1; 1 PROSITE; PS00404; UTEROGLOBIN_2; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMART; SM00096; UTG;
                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     . Similarity
26; Conser
                                                                                     Chem.
                                                                                   structure, and expression of a rat binding protein inated hiphenyls. Homology to the hormonally regulat one-binding protein uteroglobin."; Chem. 265:12690-12693(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    91 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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24
90
                                                                                                                                                                                                                                                                                                                                                                         Singh G.,
                      FROM N.A.; PubMed=2349092;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
91
24
90
9879 MW;
Katyal S.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chordata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22.2%;
                                                                                                                                                                                                                                                                                                                                                                         Brown W.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16;
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Pred. No. 0
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INTERCHAIN (WITH C-24').
587614DAE9E4820F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      UTEROGLOBIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata;
Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86
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Singh G.,
                                                                                                                                                                         Ahlgren R.,
J.;
                                                                                                                                                                                                                                                                                                                                                                       Kennedy A.L.,
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  Beato
                                                                                                                                                                                                                                                                                                            amino acid and
ssion.";
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                                                                                                                                                                                                                                                                                                                                                                     Squeglia N.,
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  Suske
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Murinae; Rattus
                                                                                                                              regulated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91;
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                                                                                                                            Query Match
Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                            PROSITE; PS00403; UTEROGLOBIN_1; 1.

PROSITE; PS00404; UTEROGLOBIN_2; 1.

PROSITE; PS00404; UTEROGLOBIN_2; 1.

PROSPONITE: PS00404; UTEROGLOBIN_2; 1.

PROSPONITE: PS00404; UTEROGLOBIN_2; 1.

PROSITE: PS00404; UTEROGLOBIN_2; 1.

PROSI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; J05536; AAA41817.1; -
EMBL; X51318; CAA35701.1; -
PIR; A36581; A36581.
PIR; S10185; S10185.
PIR; S21676; S21676.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. "There are no restrained by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entitles requires a license agreement (See http://www.isb-greenand an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR003628; Uteroglbn_sub.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
PRINTS; PR00486; UTEROGLOBIN.
ProDom; PD012475; Uteroglbn_sub; 1.
SMART; SM00096; UTG; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nat.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        + +
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Nat. Struct. Biol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRUCTURE BY NMR. MEDLINE=96069785;
                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=96069785; PubMed=7583672; Haerd T., Barnes H.J., Larsson C., "Solution structure of a mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resolution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Umland T.C., Swaminathan S., Furey W., Singh G., Pletcher J., Sax M.; "Refined structure of rat Clara cell 17 k'a protein at 3.0-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=92219263;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               X-RAY CRYSTALLOGRAPHY (3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic Acids Res. 18:2939-2946(1990)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mol. Biol. 224:441-448(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INHIBITOR OF PHOSPHOLIPASE A2.
SUBUNIT: Homodimer; antiparallel distlfide-linked.
TISSUE SPECIFICITY: CLARA CELLS (NONCILIATED CELLS
SURFACE EPITHELIUM OF THE PULMONARY LIRMAYS).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FUNCTION: BINDS PHOSPHATIDYLCHOLINE, POLYCHLORINATED BIPHENYLS (PCB) AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INDUCTION: BY GLUCOCORTICOIDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1CCD; 31-JAN-94.
1UTR; 07-DEC-95.
                                                              MKLSVCLLLVTLALCCYQANAEFCPALVSELLDI'FFISEPLFKLSLAKFDAPPEAVAAKL
MKIAITITVLMLSICCSSASSDICPGFLQVLEALLLGSESNYEAALKPFNPASDLQNAGT
                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                        96
                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                        A
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                                                                                                                                                                                                                                                           10449
                                                                                                                                                         22.2%;
                                                                                                                                                                                                                                                           X.
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                                                                                                                            Score 100; DB 1;
Pred. No. 0.00011;
2; Mismatches 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INTERCHAIN
INTERCHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLARA CELL
                                                                                                                                                                                                                                                    1A12988677B9EBEF CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3D-structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (WITH C-90'). (WITH C-24').
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PHOSPHOLIPID-BINDING PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     http://www.isb-sib.ch/announce/
                                                                                                                               41;
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                                                                                                                                                                                   Length
                                                                                                                               Indels
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in in
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UGR1_MOUSE
RESULT 11
                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                         Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                   VARSPLIC
VARSPLIC
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Signal: Alternative splicing.
SIGNAL 1 21
CHAIN 22 139
VARSPLIC 107 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; AF274959; AAL25708.1; -.
EMBL; AF274960; AAL25709.1; -.
EMBL; AF274961; AAL25710.1; -.
InterPro; IPR000329; Uteroglobin.
Pfam; PF01099; Uteroglobin; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mol.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "UGRP1, a uteroglobin/clara cell secretory a novel lung-enriched downstream target ger homeodomain transcription factor.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Uteroglobin-related protein 1 precursor (Secr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UGR1_MOUSE STANDARD;
Q920H1; Q920H2; Q920H3;
15-JUN-2002 (Rel. 41, Cr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ÷
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nilmi T., Keck-Waggoner C.L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                *1550E=Lung;
MEDLINE=21539178;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SCGB3A2 OR UGRP1.
                                                                       57
                                                                                                                       52 PPEAVAAKLGVKRCTDOMSLQKRSLIAEVLVKILKKCS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
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                                                                                                                                                                                                       1 MKLSVCLLLVTLALCCYQANA------EFCPALVSELLDFFFISEPLFKLSLAKFDA 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCELLULAR LOCATION: Secreted.
ALTERNATIVE PRODUCTS: 3 isoforms; A, B and C (shown here); are produced by alternative splicing.
TISSUE SPECIFICITY: Highly expressed in lung.
SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUBUNIT: Homodimer
                                                                       SVEHLVT -- GLKKCVDELGPEASEAVKKLLVIII--CS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QLKRLVDTLPQETRINIVKLTEKIL
                                                                                                                                                                          MKLVSIFLLVTIGICGYSATALLINRLPVVDKLPVPLDDIIPSF---DPL-KMLLKTLGI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Endocrinol. 15:2021-2036(2001).
                                                                                                                                                                                                                                                                                 . Similarity
27; Conser
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92
139
                                                                                                                                                                                                                                                                                 Conservative
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139
                                                                                                                                                                                                                                                                                                                                                                                   15431 MW;
                                                                                                                                                                                                                                                                                                    17.9%;
27.6%;
                                                                                                                                                                                                                                                                                    19;
                                                                                                                                                                                                                                                                            Score 80.5; E
Pred. No. 0.02
L9; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                      UTEROGLOBIN-RELATED PROTEIN 1.
VSVLFLEMICAYPRDSKKQTFAFIERVFEQSKL
HLV (IN ISOFORM B).
VIIICSY -> EALSHLV (IN ISOFORM A).
MISSING (IN ISOFORM A).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Popescu N.C., Zhou Y., Levitt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ory protein-related gene for the T/EBP/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Secretoglobin
                                                                                                                                                                                                                                                                                                         DB
.027;
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                                                                                                                                                                                                                                                                              17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SΑ
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                                                                                                                                                                                                                                                                                                                                                                           RESULT 12
CYTF_MOUSE
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Best Local S
Matches 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Signal.
SIGNAL
                                                                                                                                                                                             CYTF_MOUSE STANDARD; PRT; 144 AA G89098; 15-DEC-1998 (Rel. 37, Created) 15-DEC-1998 (Rel. 37, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) 15-JUN-2002 (Rel. 41, Last annotation update) Cystatin F precursor (Leukocystatin) (Cystametastasis-associated protein) (CMAP).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  a novel lung-enriched downstream target gene for the T/EBP/NKX2.1 homeodomain transcription factor."; Wol. Endocrinol. 15:2021-2036(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UGR1_HUMAN
Q96PL1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
MEDLINE-21539178; PubMed-11682631;
Niimi T., Keck-Waggoner C.L., Popescu N.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUN-2002 (Rel. 15-J
                                                                                             Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
SEQUENCE FROM N.A. MEDLINE=98298157;
                                                                                                                                              Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genew;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; AF313455; AAL26215.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- SUBCELLULAR LOCATION: Secreted.
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                                                                      NCBI_TaxID=10090;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -!- TISSUE SPECIFICITY: Highly expressed in lung.-!- SIMILARITY: BELONGS TO THE UTEROGLOBIN FAMILY. UGRP SUBFAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID-9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          50 DAPPEAVAAKLGVKRCTDQMSLQKRSLIAEVLVKILKKCS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MKLSVCLLLVTLALCCYQANA------EFCPALVSELLDFFFISEPLFKLSLAKF 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GISVEHLVE--GLRKCVNELGPE----ASEAVKKLLEALS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MKLVTIFLLVTISLCSYSATAFLINKVPLPVDKLAPLPLDNILPFM---DPL-KLLLKTL 56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
27; Conser
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41, Last sequence update)
41, Last annotation update)
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PubMed=9632704;
                                                                                             Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10161 MW;
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93
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                                                                                                                                                                                                                       (Leukocystatin) (Cystatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 75.5;
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   POTENTIAL.
UTEROGLOBIN-RELATED PROTEIN
                                                                                             Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Craniata; Vertebrata;
Catarrhini; Hominidae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FBD4BFAC2BF33718 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                    update)
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.067;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34;
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                                                                                                                                                                                                                          Cystatin-like
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                                                                                                Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Levitt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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RESULT 13
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IRA2_YI
AC P19158
AC P19158
AC P19158
AC P19258
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DISULFID
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TRA2_YEAST STANDARD; PRT; 3079 AA. P19158; Q08239; D1-NOV-1990 (Rel. 16, Created) O1-NOV-1990 (Rel. 16, Last sequence update) O1-NOV-1997 (Rel. 35, Last annotation update) Inhibitory regulator protein IRA2.

IRA2 OR GLC4 OR CCS1 OR YOLO81W OR 00985.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CHAIN
ACT_SITE
                                                                       protein with a domain homologous
protein.";
                                                                                                                    MEDLINE-90318397; PubMed-2164637;
Tanaka K., Nakafuku M., Tamanoi F., Ka
"IRA2, a second gene of Saccharomyces
                                                                                                                                                                                                                                                              Saccharomyces cerevisiae (Baker's yeast).

Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL;
HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; AF031825; AAC40140.1; EMBL; AF031825; AAC40139.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss institute of Bioinformatics and the EMBL outstation the European Bioinformatics institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                     SEQUENCE FROM N.A. MEDLINE-90318397;
                                                                                                                                                                                                                                        NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pfam; PF00031; cystatin; SMART; SM00043; CY; 1.
       SEQUENCE OF 1-2423
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                                                 10:4303-4313(1990)
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chibitor; Glycoprotein;
potenvial.
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SECONDARY AREA OF C
BY SIMILARITY.
BY SIMILARITY.
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Pred. No. 0.57;
Pred. No. 0.57;
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                                                                                                GTPase-activating
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                                                                                                                                            Toh-E A.;
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Best Local
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EMBL; X83121; CAA58201.1; -.
EMBL; X75449; CAA53202.1; -.
EMBL; Z74823; CAA99093.1; -.
PIR; S11190; RGBYIZ.
SGD; S0005441; IRA2.
InterPro; IPR001936; RasGAP.
Pfam; PE00616; RasGAP; 1.
                                                                                                                                                                CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IDENTIFICATION OF CCS1 AS IRA2.

MEDILINE-92405329; PubMed-1326414;

Bussereau F., Dupont C.H., Boy-Marcotte E., Mallet L.,

"The CCS1 gene from Saccharomyces cerevisiae which is i
mitochondrial functions is identified as IRA2 an attenu
and RAS2 gene products.";

Curr. Genet. 21:325-329(1992).

-i- FUNCTION: INHIBITORY REGULATOR OF THE RAS-CYCLIC AM
STIMULATES THE GTPASE ACTIVITY OF RAS PROTEINS.

-i- SIMILARITY: CONTAINS 1 RAS-GAP DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                            the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed. entitles requires a license agreement (; ee or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tzermia M., Katsoulou C., Alexandraki D., "Sequence analysis of a 33.2 kb segment ::rom the left arm chromosome XV reveals eight known genes and ten new open r frames including homologues of ABC transporters, inositol phosphatases and human expressed sequence tags."; yeast 13:583-589(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zumstein E., Griffin H., Schweizer M.:
Zumstein E., Griffin H., Schweizer M.:
"Sequence of a 10.27 kb segment on the
Saccharomyces cerevisiae includes part
1914
                                                     1854
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DOMAIN
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                                                                                                                                                                                           DOMAIN
                                                                                                                                                                                                                                                GTPase activation. DOMAIN 1701
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-S288c / FY1679;
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RI
                                                                               FCPALVS----ELLDFFFISEPLFKLSLAK------
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RAS_GTPASE_ACTIV_2;
                                                                                                                       14.8%;
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Pred. No. 24;
9; Mismatches
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POLY-SER.
POLY-SER.
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651EB::A2EEB479C0 CRC64;
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                                                                               -FDAPPEAVAAKLGVKRCTD 67
                                                                                                                                    Length 3079;
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SY04_CHICK
ID SY04_CHICK
AC Q90826; Q91

Q910C9

STANDARD;

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RESULT 14

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RESULT 15
FEL2_FELCA
ID FEL2_F
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P30440;
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SCYA4.
                                                                                                                                                                                                                                                               DISULFID DISULFID
                                                                                                                                                                                                                                                                                                                                          InterPro; IPR000827; CC_chemkine_sml.
InterPro; IPR001811; Chemokine_IL8.
Pfam; PF00048; IL8; 1.
SMART; SM00199; SCY; 1.
SMART; SM00199; SCY; 1.
CHECKER, PS00472; SMALL_CYTOKINES_CC; CHECKER, PS00472; CHECK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the European Bloinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE=95369710; PubMed=7642115; Petrenko O., Ischenko I., Enrietto P.J.; Petrenko O., Ischenko I., Enrietto P.J.; Chicken chemokine "Isolation of a cDNA encoding a novel chicken chemokine mammalian macrophage inflammatory protein-1 beta."; Gene 160:305-306(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-NOV-1997 (Rel. 35, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Small inducible cytokine A4 homolog precursor
                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; L34553; AAA48747.1; -.
EMBL; AJ243034; CAB45103.1; -.
HSSP; P13236; 1HUM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hughes S.M., Bumstead "Mapping of the gene chemokine SCYA4.";
                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (JUL-1994) to the -!- FUNCTION: MONOKINE WITH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gallus gallus (Chicken).
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                                                                                                   CONFLICT
                                                                                                                                                                                                                                                                                                                                Cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        between
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SUBUNIT: HOMODIMER (BY SIMILARITY).
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                                                                                                                                   MKLSVCLLLVTLALCCYQANA-
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                                                                                                                                                                                  Similarity
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UL-1994) to the
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                     STANDARD;
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SMALL INDUCIBLE CYTOKINE A
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50AF9679A267408F CRC64;
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; Galliformes; Phasianidae; Phasianinae;
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CHEMOKINETIC PROPERTIES
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01-APR-1993 (Re
01-APR-1993 (Re
15-JUN-2002 (Re
Major allergen
                                                                                                                                             CARBOHYD
VARSPLIC
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"Cat allergen 1: Biochemical, antigenic, and allergenic properties.";

J. Allergy Clin. Immunol. 74:147-153(1984).

-I- SUBUNIT: HETEROTETRAMER COMPOSED OF TWO NON-COVALENTLY LINKED DISULFIDE-LINKED HETERODIMER OF CHAINS 1 AND 2.

-I- ALTERNATIVE PRODUCTS: 3 isoforms; a long form/CH2L (shown here), a short form/CH2S and a truncated form/CH2ST; may be produced by alternative splicing.

-I- TISSUE SPECIFICITY: THE LONG FORM IS PREFERENTIALLY EXPRESSED IN THE SALIVARY GLAND, WHILE THE SHORT FORM IS PREFERENTIALLY
                                                                                                                                                                                                                  SIGNAL
                                                                                                                                                                                                                                                                                                              EMBL; M77341; AAC41616.1; EMBL; X62478; CAA44345.1;
                                                                                                                                                                                                                                                                                                                                                                                                                         between the Swiss Institute of Bioinformatics and the E
the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by an
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MEDLINE=84265679; PubMed=6747135;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE OF 18-37, AND CHARACTERIZAT MEDLINE-91287714; PubMed-1712068; Duffort O.A., Carreira J., Nitti G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             major allergen from the domestic Gene 113:263-268(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A., AND SEQUENCE OF 18-100.
MEDLINE-92052157; PubMed-1946388;
Morgenstern J.P., Griffith I.J., Brauer A.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Carnivora; Fissipedia; Felidae; Felis.
                                                                                                                                                                                                                                                                                                                                                                                 or send an email to license@isb-sib.
                                                                                                                                                                                                                                                                                                                                                                                               modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          domesticus
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"Amino acid sequence of fel dI, the major allergen
cat: protein sequence analysis and cDNA cloning.";
Proc. Natl. Acad. Sci. U.S.A. 88:9690-9694(1991).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Fel d I-B)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DISEASE: MAJOR ALLERGEN PRODUCED BY THE DOMESTIC CAT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EXPRESSED IN THE SKIN.
                                                                                                                                                                                                                                                                JC1127; JC1127
JC1145; JC1145
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(Rel. 25, Last sequence update)
(Rel. 41, Last annotation update)
gen I polypeptide chain 2 precursor (Allergen Fel d 1-B)
(Allergen Cat-1) (AG4) (FdI).
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QFRTP (IN ISOFORM CH2ST).
TTISSSKD -> IAINEY (IN ISOFORM
I -> L (IN CH2LV).
I -> V (IN CH2LV).
RV -> KF (IN CH2SV).
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major allergen
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A;Residues: 1-111 <PAR>
A;Residues: 1-111 <PAR>
R;Liao, S; Chen, C; Huang, I.Y.
J Biol. Chem. 257, 122-125, 1982
A;Title: Prostate alpha-protein. Complete arino acid sequence of the component that i A;Title: Prostate alpha-protein. Complete arino acid sequence of the component that i A;Title: Prostate alpha-protein. Complete arino acid sequence of the component that i A;Title: Prostate alpha-protein. Complete arino acid sequence of the component that i A;Title: Prostate alpha-protein. A;Title: Identification of a complementary-iNA clone containing part of the sequence A;Reference number: A90348
A;Accession: A90348
A;Accession: A90348
A;Accession: A90348
A;Molecule type: mRNA
A;Residues: 13-14, 'S',16, 'GG',19-65 <DEL>
R;Delaey, B.; Dirckx, L.; Peeters, B.; Volc(aert, G.; Mous, J.; Heyns, W.; Rombauts, Eur. J. Biochem. 133, 645-649, 1983
A;Title: The nucleotide sequence of cDNA complementary to the C(1) component of rat [A;Reference number: S42615; MUID:83234456; PMID:6688048
A;Accession: S42615
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F;1-23/Domain: signal sequence #status precicted <SIG>
F;24-111/Product: prostatic steroid-binding protein chain Cl #status experimental
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 15-Oct-1982 #sequence_revision 15-Oct-1982 #text_change 22-Jun-1999
C;Accession: A93286; A92348; A90348; S42615; A03252
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Best Local Similarity
60 LGVKRCTDQMSLQKRSLIAEVLVKILKKCSV 90
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                                                                                     VELSICILIM-LAVCCYEANASQICELVAHET::SFLMKSEEELKKELEMYNAPPAAVEAK 62
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A;Title: Prostatic steroid binding protein: gene duplication and steroid bind A;Reference number: A93286; MUID:82220075; PMID:6896362
A;Recession: A03251
A;Accession: A03251
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A;Accession: A03261
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A;Accession: A03261
A;Cross-references: GB:000776; NID:9206448; PIDN:AAA51641.1; PID:9206450
A;Cross-references: GB:000776; NID:9206448; PIDN:AAA51641.1; PID:9206450
A;Cross-references: A26671; Decourt, J.L.; Claessens, F.; Peeters, B.; Rombauts
Nucleic Acids Res. 15, 1627-1641, 1987
A;Cross-reference number: A26671; MUID:87146484; PMID:2881277
A;Accession: A26671
A;Molecule type: DNA
A;Residues: 1-25, 'Q', 26-86, 'I', 88-94, 'VWLQINFPRGRWFSEIN' <DEL>
A;Cross-references: GB:X05034; NID:956857; PIDN:CAA28708.1; PID:956858
C;Comment: Steroid-binding protein, the principal secretory protein in rat pi
C;Roywords: heterotetamer; prostate; steroid binding
F;1-20/Domain: signal sequence #status predicted <SIG>
F;21-98/Product: prostatic steroid-binding protein chain C2 #status predicted
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A;Title: Cloning of a SA;Reference number: S68
A;Accession: S68231
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  prostatic steroid-binding protein chain C2 precursor - C;Species: Rattus norvegicus (Norway rat) C;Date: 15-Oct-1982 #sequence_revision 15-Oct-1982 #tex C;Accession: A03251; A26671 R;Parker, M.; Needham, M.; White, R.
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C;Superfamily: uteroglobin
F;1-21/Domain: signal sequence |
F;22-95/Product: FHG22 protein |
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C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 21-Aug-1998
C;Accession: S68231
R;Dominguez, P.
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A; Residues: 1-95 < DOM>
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|1-21/Domain: signal sequence #status predicted <SIG>
|2-95/Product: FHG22 protein #status predicted <MAT>
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                                                 MRLSLCLLTI-LVVCCYEANGQTLAGVCQALQDVTITFLLNPEEELKRELEEFDAPPEAV 59
                                                                                                   MKLSVCLLLVTLALCCYQAN----AEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAV 56
     AAKLGVKRCTDQMSLQKRSLIAEVLVKILKKCSV
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g of a Syrian hamster cDNA :
ber: S68231; MUID:96105393;
                                                                                                                                                                                                                                                       prostatic steroid-binding protein chain C2 #status predicted
                                                                                                                                                         Conservative
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                                                                                                                                                  Score 166.5; DB 1;
Pred. No. 4.6e-12;
Pred. No. 4.6e-12;
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; PMID:7498554
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A; Nolecule type: protein
A; Residues: 22-23,'X',25-28,'X',30-31,'X',33-36 <SIZ>
A; Residues: 22-23,'X',25-28,'X',30-31,'X',33-36 <SIZ>
A; Renard, A.; Roels, H.; Lauwerys, R.; Witters, R.; Gielens, Clin. Chim. Acta 207, 239-249, 1992
Clin. Chim. Acta 207, 239-249, 1992
A; Title: Human urinary protein 1: evidence for identity with
                    C:Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 28-Feb-1980 #sequence_revision 15-Oct-1982 #text_change 15-Sep-2000
C;Accession: A92391; A93461; A90935; A24217; A90303; A90417; A93824; A94608;
R;Bailly, A.; Atger, M.; Atger, P.; Cerbon, M.A.; Alizon, M.; Vu Hai, M.T.;
J. Biol. Chem. 258, 10384-10389, 1983
A;Title: The rabbit uteroglobin gene. Structure and interaction with the profix Reference number: A92391; MUID:83290960; PMID:6309802
A;Accession: A92391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C;Superfamily: uteroglobin
F;1-21/Domain: signal sequence #status predicted <SIG>
F;22-91/Product: Clara cell 10K protein #status experi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A:Experimental source: urine A;Note: sequence extracted from NCBI backbone (NCBIP:119391) R;Hay, J.G.; Danel, C.; Chu, C.; Crystal, R.G. Am. J. Physiol. 268, 565-575, 1995
A;Title: Human CC10 gene expression in airway epithelium and A;Reference number: I38397
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A:Title: Amino-acid and cDNA nucleotide sequences of human Clara cell 10kDa protein.
A:Reference number: JS0036; MUID:89000784; PMID:3167058
                                                                                                                                                                                                                             uteroglobin precursor [validated] - rabbit N; Alternate names: blastokinin
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C; Date: 31-
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A; Residues: 1-91 <RES>
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A;Accession: PS0309
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A; Residues: 1-91 <SIN>
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31-Mar-1992 #sequence
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R;Atger, M.; Mercler, J.C.; Haze, G.; Fridlansky, F.; Milgrom, E.
Blochem. J. 177, 985-988, 1979
A;Title: M. terminal sequences of uteroglobin and its precursor.
A;Reference number: A90303; MUID:79187160; PMID:571719
A;Accession: A90303
A;Molecule type: protein
A;Residues: 1-5, 'F',7-10, 'X',15,'G',17-54,'X',56,'B',58-66,'B',68-70,'XX',73 <ATG>
R;Ponstingl, H.; Nieto, A.; Beato, M.
Blochemistry 17, 3908-3912, 1978
A;Fitle: Amino acid sequence of progesterone-induced rabbit uteroglobin.
A;Recession: A90417; MUID:79042086; PMID:568483
A;Accession: A90417; MUID:79042086; PMID:568483
A;Recession: A90417; MUID:79042086; PMID:568483
A;Residues: 22-81,'Q',83-91 < PON>
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A; Residues: 50-62;67-71 <PO2>
R: Morize, I.; Surcouf, E.; Vaney, M.C.; Buehner, M.; Mornon,
submitted to the Brookhaven Protein Data Bank, April 1989
A; Reference number: A50025; PDB:10TG
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A;Title: Amino acid sequence of a progesterone-binding pro A;Reference number: A93824; MUID:79074850; PMID:281700
A;Accession: A93824
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A;Cross-references: GB:K00049; NID:g165789
R;Suske, G.; Wenz, M.; Cato, A.C.B.; Beato, M.
Nucleic Acids Res. 11, 2257-2271, 1983
N;Title: The uteroglobin gene region: hormonal regulation,
A;Reference number: A93461; MUID:83220783; PMID:6304644
A;Accession: A93461
A;Title: Structure and refinement of the oxidized P2-1 form of uteroglobin at 1.64 angst A;Reference number: A44653; MUID:8919637; PMID:2704039
A;Contents: annotation; X:ray crystallography, 1.64 angstroms; disulfide bonds R;Menne, C.; Suske, G.; Arnemann, J.; Wenz, M.; Cato, A.C.B.; Beato, M.
                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Contents: annotation; X-ray crystallography, 1.34 angstroms, residues 22-91 R;Morize, I.; Surcouf, E.; Vaney, M.C.; Epelboin, Y.; Buehner, M.; Fridlansky, J. Mol. Biol. 194, 725-739, 1987
A;Title: Refinement of the C222-1 crystal form of oxidized uteroglobin at 1.34
A;Reference number: A44652; MUID:88011213; PMID:3656405
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A; Residues: 22-49, 'D,'51, 'EN',54-59,61-66, 'NEPSL',72-91 <POP>
R; Popp, R.A.; Foresman, K.R.; Wise, L.D.; Daniel Jr., J.C.
submitted to the Atlas, October 1982
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                                                                                                                                                              A;Contents: annotation; X-ray crystallography, 1.64 angstroms, residues R;Bally, R.; Delettre, J. J. Mol. Biol. 206, 153-170, 1989
                                                                                                                                                                                                                                                                                                                                                     A; Contents: annotation; X-ray crystallography, 1.34 R; Bally, R.; Delettre, J.
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A; Accession: A94608
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A; Residues: 22-91 <LOP>
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A; Residues: 1-91 <CHA>
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A; Residues: 1-91 <SUS>
                                                                                                                                                                                                                                                                                submitted to the Brookhaven Protein Data Bank,
A; Reference number: A50553; PDB: 2UTG
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A;Cross-references: GB:M25057; NID:g165802; PiDN:AAA31498.1; PID:g16R;Suske, G.; Menne, C.; Cato, A.; Wenz, M.; B:Mato, M. Prog. Clin. Biol. Res. 85, 139-146, 1982
A;Title: Structure and regulated expression of the uteroglobin gene A;Reference number: I46907; MUID:82275176; PM:D:6287481
A;Accession: I46907
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Biochem. Biophys. Res. Commun. 95, 197-204, 1380
A;Title: Cloning of the rabbit uteroglobin st:uctural gene.
A;Reference number: 146905; MUID:81021016; PM(D:7417250)
A;Accession: 146905
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C;Complex: homodimer linked by two disulfide konds
C;Superfamily: uteroglobin
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A; Residues: 53-66,'NT', 69-72 <AT2>
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A;Residues: 1-45,'V',47-91 <SU2>
A;Residues: 1-45,'V',47-91 <SU2>
A;Cross-references: GB:M32012; NID:g165807; P:DN:AAA31500.1; PID:g165809
R;Atger, M.; Perricaudet, M.; Tiollais, P.; M.:lgrom, E.
Biochem. Biophys. Res. Commun. 93, 1082-1088, 1980
A;Title: Bacterial cloning of the rabbit utervglobin structural gene.
A;Reference number: 146906; MUID:80241888; PM:D:615676
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A;Title: Isolation and structure of the gene for the progesterone-inducible protein A;Reference number: 146904; MUID:83014990; PMID:6956897
A;Accession: 146904
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A; Residues: 39-77 <CH2>
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61 QMKKVLDSLPQTTRENIMKLTEKIVK
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A;Reference number: A53025; MUID:94292183; PMID:8020953
A;Accession: A53025
A;Status: preliminary

R;Stripp, B.R.; Huffman, J.A.; Bohinski, R.J. Genomics 20, 27-35, 1994
A;Title: Structure and regulation of the muri

C;Species: Mus musculus (house mouse)
C;Date: 03-May-1994 #sequence_revision 21-Jan-1997
C;Accession: A53025; A56656; I51925; S24783

#text_change 22-Jun-1999

secretory protein

gene

uteroglobin precursor - mouse N; Alternate names: CC10; Clara cell 10K protein precursor; Clara cell secretory

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A36581
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N;Alternate names: Clara cell 10k secretory protein
N;Alternate names: Clara cell 10k secretory protein
C;Specles: Rattus norvegicus (Norway rat)
C;Date: 08-Mar-1991 #sequence_revision 08-Mar-1991 #text_change 20-Aug-1999
C;Accession: A36581; S10185; S21676
R;Nordlund-Moeller, L.; Andersson, O.; Ahlgren, R.; Schilling, J.; Gillner, M.; Gustafss
J, Blol. Chem. 265, 12690-12693, 1990
A;Title: Cloning, structure, and expression of a rat binding protein for polychlorinated
A;Accession: A36581; MUID:90324266; PMID:2115524
A;Accession: A36581; MUID:90324266; PMID:2115524
A;Accession: A9684
A;Residues: 1-96 <NDS;
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-96 <NDS;
A;Cross-references: GB:J05536; NID:9206039; PIDN:AAA41817.1; PID:9206040
R;Hagen, G; Wolf, M.; Katyal, S, L.; Singth, G; Beato, M.; Suske, G.
Nucleic Acids Res. 18, 2939-2946, 1990
A;Title: Tissue-specific expression, hormonal regulation and 5'-flanking gene region of
A;Accession: S10185
A;Accession: S10185; MUID:90272398; PMID:2349092
A;Status: translation not shown
A;Residues: 1-18 <HAGS
A;Cross-references: EMBL.X51318; NID:955336; PIDN:CAA35701.1; PID:935537
R;Umland, T.C.; Swaminathan, S.; Furey, W.; Singh, G.; Pletcher, J.; Sax, M.
J. Mol. Btol. 224, 441-448, 1992
A;Title: Refined structure of rat Clara cell 17 kDa protein at 3.0 A resolution.
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A; Note: sequence extracted from NCBI backbone (NCBIP:126148)
A; Note: parts of this sequence, including the amino end of the m
R; Margraf, L.R.; Finegold, M.J.; Stanley, L.A.; Major, A.; Nawki
Am. J. Respir. Cell Mol. Biol. 9, 231-238, 1993
A; Title: Cloning and tissue-specific expression of the cDNA for A; Reference number: 151925; MUID:94000840; PMID:8398159
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A;Cross·references: GB.L24372; NID:g461147; PIDN:AAA65446.1; PID:g785054
R;Singh, G.; Katyal, S.L.; Brown, W.E.; Kennedy, A.L.
Exp. Lung Res. 19, 67-75, 1993
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Pred. No. 0.00018;
9; Mismatches 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the mouse Clara
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         A; Molecule type: DNA
A; Residues: 1-113 <RAY>
C; Comment: This protein is the major secretory product of the Clara
C; Superfamily: uteroglobin
F;73/Region: ochre stop codon
                                                                                                                                                                                                                                                                              R;Ray, M.K.; Magdaleno, S.; O'Malley, B.W.; DeMayo, F.J. Blochem. Blophys. Res. Commun. 197, 163-171, 1993
A;Title: Cloning and characterization of the mouse clara A;Reference number: JC2026; MUID:94071937; PMID:7916613
                                                                                                                                                                                                                                                                                                                                                                                                                                                          cell specific 10K protein - mouse C; Species: Mus musculus (house mouse) C; Date: 27-Jun-1994 #sequence_revision C; Accession: JC2026
                                                                                                                                                                                                                                    A; Reference number: A; Accession: JC2026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           δÃ
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61

61 GVKRCTDQMSLQKRSLIAEVLVKILK

86 98

27-Jun-1994 #text_change

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10

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protein

cell and

binds

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C;Superfamily: uteroglobin
C;Keywords: lung; steroid binding; uterus
F;1-21/Domain: signal sequence *status predicted <SIG>
F;22-91/Product: uteroglobin *status predicted <MAT>
F;24/Disulfide bonds: interchain (to 90) *status predicted
F;90/Disulfide bonds: interchain (to 24) *status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Species: Lepus capensis (brown hare)
C;Date: 30-Sep-1987 #sequence_revision 30-Sep-1987 #text_change 22-Jun-1999
C;Accession: A23825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A:Reference number: S21676; MUID:92219263; PMID:1560460 A;Contents: annotation, X-ray crystallography, 3.0 angstroms C;Superfamily: uteroglobin F;24/Disulfide bonds: interchain (to 90) #status experimenta F;90/Disulfide bonds: interchain (to 24) #status experimenta
                                                                                                                                                                                                                                                                                                       C;Comment: Uteroglobin, synthesized in the uterus and lung, is secreted C;Complex: homodimer linked by two disulfide bonds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Lopez de Haro,
                                                                                                                                                                                                                                                                                                                                                                                                  A; Molecule type: mRNA
A; Residues: 1-91 <LOP>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Title: Nucleotide and derived amino acid sequences of a A; Reference number: A23825; MUID:86323069; PMID:3019311
                                                                                                                                                                                                                                                                                                                                                        A; Experimental source: lung
                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: GB:M25609; NID:g164246; PIDN:AAA30960.1; PID:g164247
                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Accession: A23825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N; Alternate
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                                                                                                           Local
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                                         1 MKLSVCLLLVTLALCCYQANAEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAAKL 60
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MKLTITLALVTLALLCSPASAGICPGFAHVIENLLLGTPSSYGTSLKEFQPDDAMKDAGM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             names: blastokinin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  precursor - brown hare
                                                                                                           Similarity
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                                                                                        Conservative
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                                                                                                      21.6%;
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                                                                                   Score 97; DB
Pred. No. 0.00
15; Mismatches
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Pred. No. 0.00023;
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#status experimental
                                                                                                                                  DB 1;
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major allergen chain 2 precursor, short form - cat
C;Species: Felis silvestris catus (domestic cat)
C;Decies: Felis silvestris catus (domestic cat)
C;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 05-Nov-1999
C;Accession: JC1127
R;Griffith, I.J.; Craig, S.; Pollock, J.; Yu, X.B.; Morgenstern, J.P.; Rogers,
Gene 113, 263-268, 1992
A;Title: Expression and genomic structure of the genes encoding FdI, the major
A;Reference number: JC1126; MUID:92241678; PMID:1572548
A;Accession: JC1127
A;Molecule type: DNA
A;Residues: 1-107 GGRI>
A;Cross-references: GB:X62478; NID:g395406; PIDN:CAA44345.1; PID:g395407
A;Experimental source: skin
C;Genetics:
                                                                                                                                                           A;Introns: 21/1; 81/3
C;Reywords: glycoprotein
F;1-17/Demain: signal sequence #status predicted <SIG>
F;18-107/Product: major allergen chain 2, short form #status predicted
F;50/Binding site: carbohydrate (Asn) (covalent) #status predicted
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A;Residues: 1-94 <DEA>
A;Cross-references: EMBL:X60661; NID:g57735; PIDN:CAA43068.1; PID:g57736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBO J. 10, 2813-2819, 1991

A; Title: Novel genes for potential ligand-binding proteins A; Reference number: S17447; MUID:92007724; PMID:1915264
A; Accession: S17449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Rattus norvegicus (Norway rat)
C;Date: 13-Jan-1995 #sequence_revision 13
C;Accession: S17449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     probable ligand-binding protein RYD5 - rat
C; Species: Rattus norvegicus (Norway rat)
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                               δ
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                                                                                                                                                                                                                                                            A;Gene: Ch2
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Best Local
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Best Local
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                                                                                                                                                         /Binding site: carbohydrate (Asn) (covalent) #status
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MKLSVCLLLVTLALC-CYQANAEFCPALVSELLDFFFISEP--LFKLSLAKFDAPPEAVA 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASLKPFNPGSDLXNAGTQLKRLVDTLPQETRINIMKLTEKIL 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MKIAITITVVMLSICCSSLLEKKLLSFCFFLPATPASSDICPGFLQVLEALLMESESGYV 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MKLSVCLLLVTLALCCYQ------ANAEFCPALVSELLDFFFISEPLFK 43
                                     LSVCLLLVTLALCCYQANAEFCPALVSELLDFFFI----SEPLFKLSLAKFDA-PPEAVA 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AKLGVKRCTDQMSLQKRSLIAEVLVKIL 85
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  LLVLALLVTQALGVKM--AETCPI----FYDVFFAVANGNELLLDLSLTKVNATEPERTA 58
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25; Conser
                                                                                               Similarity
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                                                                              Conservative
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28.4%;
                                                                                               15.6%;
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23.5%;
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Pred. No. 0.022;
                                                                          Score 70; DB 2;
Pred. No. 0.76;
B; Mismatches 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                              38;
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                                                                                                                 Length 107
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                                                                            12;
                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                   the major allergen
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                                                                                                                                                                             <MAT>
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T01369
                                                                                                                                                                         A; Reference number: Z19930
A; Accession: T24735
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A;Residues: 1-1623 <STO>
A;Cross-references: GB:A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Accession: T01369
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A; Map position:
A; Introns: 42/2
                                                            C; Genetics:
                                                                                                                   A; Molecule type: DNA
A; Residues: 1-578 <WIL>
                                                                                                                                        A; Molecule
                                                                                                                                                                                                                                     R; McLay,
                                                                                                                                                                                                                                                      C; Accession:
                                                                                                                                                                                                                                                                                                                                                           RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Gene: At2g34660; T29F13.13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C;Genetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Status: preliminary
                                     A; Gene: CESP:T09E11.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                     104
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  47
                                                                                                                                                                                                                                                                                                                                                                                                                                                          52 P--PEAVAAKLGVK 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
                                                                                                                                                                                                                                                                                                                                                                                                                     PGLPPFEAFGLGVK 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AKLGVKRCTDQMSLQKRSLIAEVLVKILKKC
  42/2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15.3%;
1 Similarity 36.5%;
27; Conservative
177/1; 257/2; 317/1; 352/3; 393/1; 430/2; 497/3
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A;Cross-references: EMBL:AC003096; NID:g3132469; PID:g3132479
A;Experimental source: cultivar Columbia
R;Lin, X.; Kaul, S.; Rounsley, S.D.; Shea, T.F.; Benito, M.I.; Town, C.D.; Fujii, C.X M.; Koo, H.; Moffat, K.S.; Cronin, L.A.; Shen, M.; VanAken, S.E.; Umayam, L.; Tallon, euss, D.; Nlerman, W.C.; White, O.; Eisen, J.A.; Salzberg, S.L.; Fraser, C.M.; Venter, Nature 402, 761-768, 1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R;Rounsley S.D.; Lin, X.; Ketchum, K.A.; Crost submitted to the EMEL Data Library, May 1998 A;Description: Arabidopsis thaliana chromosome A;Reference number: Z14179
                                                                                                                                                                                                                                                                                                                                             hypothetical protein T09E11.5 - Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Species: Caenorhabditis elegans C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 06-Oct-2000 C;Accession: T24735
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Map position: 2
A;Introns: 110/3; 170/2; 183/3; 236/1; 274/2; 301/2; 309/3; 328/3; 350/3; 375/3; 411/C;Superfamily: human multidrug resistance protiain cMOAT2; ATP-binding cassette homolo F;1259-1452/Domain: ATP-binding cassette homology <ABC>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Title: Sequence and analysis of chromosome 2 of the plant Arabidopsis thaliana A;Reference number: A84420; MUID:20083487; PMID:10617197 A;Accession: D84759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-1623 <ROU>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Date: 12-Feb-1999 #sequence_revision 12-Feb·1999 #text_change 23-Mar-2001
C;Accession: T01369; D84759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABC transporter AtMRP2 [imported] - Arabidops:s thaliana N;Alternate names: multidrug-resistance protein homolog C;Species: Arabidopsis thaliana (mouse-ear cress)
A;Cross-references: EMBL:Z81147; PIDN:CAB03529.1; GSPDB:GN00019; CESP:T09E11.5
A;Experimental source: clone T09E11
                                                                                                                                                A; Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                 submitted to the EMBL Data Library, October 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3 LSVCLLLVTLALCCYQANAEFCPALVSELLDFFF----ISEPLFKL----SLAKFDA 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LVLCLYRIWLALKDHKVE-RFC--LRSRLYNYFLALIAAYATAEPLFRLIMGISVLDFDG 103
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GB:AE002093; NID:g3132479; PIDN:AAC16268.1; GSPDB:GN00139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 69; DI
Pred. No. 14,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DN 2; Length 1623;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ΙΙ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22;
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R.Zumstein, E.; Griffin, H.; Schweizer, M.

Yeast 10, 1383-1387, 1994

A.; Ittle: Sequence of a 10.27 kb segment on the left arm of chromosome XV from Saccharomy
A.; Reference number: $48254

A.; Status: nucleic acid sequence not shown; translation not shown
A.; Molecule type: DNA
A.; Residues: 1-2423 <ZU2>
A.; Cross-references: EMBL:X75449; NID:g414079; PIDN:CAA53202.1; PID:g414081
A.; Onto: the nucleotide sequence was submitted to the EMBL Data Library, October 1993
R.; Zumstein, E.; Pearson, B.M.; Kalogeropoulos, A.; Schweizer, M.
A.; Reference number: $50410
A.; Accession: $50426
A.; Molecule type: DNA
A.; Residues: 1-2423 <ZU3>
A.; Cross-references: EMBL:X83121; NID:g600461; PIDN:CAA58201.1; PID:g600480
F;693-709/Domain: transmembrane #status predicted <TM1>F;1135-1151/Domain: transmembrane #status predicted <TM2>F;1701-1910/Domain: ras-specific GAP catalytic domain hom
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R; Alexandraki, D.; Katsoulou, C.; Tzermia, M.
submitted to the Protein Sequence Database, July 1996

A; Reference number: $66756

A; Recession: $66774

A; Molecule type: DNA
A; Residues: 1983-3079 <ALE>
A; Cross-references: EMBL: Z74823; GSPDB:GN00015; MIPS:YOL081w
A; Cross-references: EMBL: Z74823; GSPDB:GN00015; MIPS:YOL081w
A; Tanaka, K.; Nakafuku, M.; Tamanoli, F.; Kaziro, Y.; Matsumoto, K.; Toh-e, Mol. Cell. Biol. 10, 4303-4313, 1990
                                                                                                                                                                                                  C; Superfamily: regulatory protein IRA2; ras-specific GAP
                                                                                                                                                                                                                                                                 A; Map position: 15L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mol. Cell. Biol. 10, 4303-4313, 1990
A;Title: IRA2, a second gene of Saccharomyces cerevisiae that encodes a protein with A;Reference number: A35656; MUID:90318397; PMID:2164637
A;Accession: A35656
A;Status: nucleic acid sequence not shown
A;Molecule type: DM: Molecule type: DM: A;Molecule type: DM: A;
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C;Date: 30-Sep-1991 #sequence_revision 16-Aug-1996 #text_change 03-Dec-1999 C;Accession: S66775; S66774; A35656; S48254; S50426; S11190; S38505 R;Zumstein, E.; Pearson, B.M.; Kalogeropoulos, A.; Schweizer, M. submitted to the Protein Sequence Database, July 1996
A;Reference number: S66775
                                                                                                                                                                                                                                                                                                                 A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                A; Gene: SGD: IRA2; MIPS: YOL081w
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A;Residues: 1-2423 <ZUM>
A;Cross-references: EMBL:Z74823; GSPDB:GN00015; MIPS:YOL081w
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A; Molecule type: DN
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                                                                                                                                                             transmembrane protein
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Pred. No. 10
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domain homology <GAP>
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                                                                                                                                                                                                  catalytic domain homology
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Search completed: January Job time : 19 secs

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2003, 14:56:02

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C;Genetics:
A;Gene: AGR_C_2384
A;Map position: ci
                                                                                                                                                                                                                                                                                                                                              R;Goodner, B.; Hinkle, G.; Gattung, S.;
A.; Liu, F.; Wollam, C.; Allinger, M.;
Science 294, 2323-2328, 2001
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                                    δõ
                                                                                                                                                                                                                                                                                                                                                                                                          hypothetical protein AGR_C_2384 [imported] - Agrobacterium tumefaciens (strain C58, C ;Species: Agrobacterium tumefaciens C:Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 11-Jan-2002 C:Date: 30-Sep-2001 #sequence_revision 30-Sep-2001 #text_change 11-Jan-2002 C:Accession: G97516
                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-102 <KUR>
                                                                                                                                                                                                                                                                                                            A; Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrobacterium A; Reference number: A97359; PMID:11743194
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                                                                                                                                                                                                            A; Cross-references:
                                                                                                                                                                                                                                                                     A; Status: preliminary
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    37
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                                  12 LALCCYQAN----AEFCPALVSELLDFFFISEPLFKLSLAKFDAPPEAVAA
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LALSGLQANMLRQAVNDPGFLAGLTDFLMSHEPDLMAFCAATDTPPETVAA
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                                                                                           14.7%;
41.2%;
                                                                                                                                                     chromosome
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Pred. No.
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Pred. No. 5
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                                                                           Mismatches
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                                                                                                              DB 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ----FDAPPEAVAAKLGVKRCTD
                                                                           24;
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87
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Markelz,
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